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Additional Information



Fluorescence polarization immunoassay for rapid screening of the pesticides thiabendazole and tetraconazole in wheat

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Keywords:	fungicides, thiabendazole, tetraconazole, fluorescence polarization immunoassay, wheat

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3 **1 Fluorescence polarization immunoassay for rapid screening of the pesticides thiabendazole**
4 **and tetraconazole in wheat**
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37

38 16 **Abstract**
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40 17 Fluorescence polarization immunoassays (FPIAs) for thiabendazole and tetraconazole were first
41
42 18 developed. Tracers for FPIAs of thiabendazole and tetraconazole were synthesized and the
43
44 19 tracers’ structures were confirmed by HPLC-MS/MS. The 4-aminomethylfluorescein-labeled
45
46 20 tracers allowed achieving the best assay sensitivity and minimum reagent consumption in
47
48 21 comparison with aminofluorescein-labeled and alkyldiaminefluoresceinthiocarbamyl-labeled
49
50 22 tracers. Measurements of fluorescence polarization were performed using a portable device. The
51
52 23 developed FPIA methods were applied for the analysis of wheat. Fast and simple sample
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24 preparation technique earlier developed by authors for pesticides was adapted for thiabendazole
25 and tetraconazole. The limits of detection of thiabendazole and tetraconazole in wheat were 20
26 and 200 $\mu\text{g}/\text{kg}$, and the lower limits of quantification were 40 and 600 $\mu\text{g}/\text{kg}$, respectively. The
27 recovery test was performed by two methods—FPIA and HPLC-MS/MS. The results obtained
28 by FPIA correlated well with those obtained by HPLC-MS/MS ($r^2 = 0.9985$ for thiabendazole,
29 $r^2 = 0.9952$ for tetraconazole). Average recoveries of thiabendazole and tetraconazole were
30 74 \pm 4% and 72 \pm 3% by FPIA, and average recoveries of thiabendazole and tetraconazole were
31 86 \pm 2% and 74 \pm 1% by HPLC-MS/MS ($n=15$).

33 **Keywords:** fungicides, thiabendazole, tetraconazole, fluorescence polarization immunoassay,
34 wheat.

36 Abbreviations

37 AF – aminofluorescein;

38 AMF – 4-aminomethylfluorescein;

39 BSA – bovine serum albumin;

40 $(\text{CH}_2)_4\text{DF}$ – butylenediaminefluoresceinthiocarbamyl;

41 $(\text{CH}_2)_6\text{DF}$ – hexamethylenediaminefluoresceinthiocarbamyl;

42 CR – cross-reactivity;

43 DCC – dicyclohexylcarbodiimide;

44 EDF – ethylenediaminefluoresceinthiocarbamyl;

45 ELISA – enzyme-linked immunosorbent assay;

46 FI – fluorescence intensity;

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3 47 FITC – fluorescein isothiocyanate isomer I;
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5 48 FPIA - fluorescence polarization immunoassay;
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7 49 HPLC-MS/MS – high-performance liquid chromatography coupled with tandem mass
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9 spectrometry;
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11 50
12 51 LOD – limit of detection;
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14 52 MRL – maximum residue level;
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16 53 NHS – N-hydroxysuccinimide;
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18 54 SD – standard deviation;
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20 55 TLC – thin-layer chromatography.
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26 57 **Introduction**

28 58 Modern agriculture is an extensive industry where profit is affected by many factors,
29
30 including plant pests and diseases. Pesticides are chemical agents targeted to be toxic to living
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32 organisms that are used to minimize yield losses. Pesticide residues in foodstuffs are controlled
33
34 by legislation in many countries because of unintended pesticide toxicity for nontargeted living
35
36 organisms, including humans.
37
38

40 63 Thiabendazole and tetraconazole are fungicides with protective and curative actions used
41
42 against pathogens of fruits, vegetables, and cereals. Thiabendazole belongs to the benzimidazole
43
44 class of pesticides, and it is applied for post-harvest treatment of food crops before dispatching
45
46 for storage. Tetraconazole is a triazole class pesticide used primarily to control diseases of the
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48 vegetative organs of plants. These compounds have low toxicity to mammals [1,2]. Nevertheless,
49
50 it is necessary to control their content in foodstuffs to avoid cases of chronic poisoning. For most
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52 plant products, the Maximum Residue Levels (MRLs) of thiabendazole are fixed at 0.01–0.05
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3 70 mg/kg (lower limits of analytical determination) according to European regulations [3] and at
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5 71 0.2–5.0 mg/kg according to Russian Hygienic standards [4]. The MRLs of tetraconazole for most
6
7 72 plant products range from 0.02 to 0.3 mg/kg according to the European Union (EU) [5]. In the
8
9 73 Russian Hygienic standards, the MRLs of tetraconazole are established for cereals at 0.2 mg/kg
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11 74 and sugar beets at 0.05 mg/kg [4].
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14 75 Actually liquid and gas chromatography with different types of detectors are the main tools
15
16 76 for pesticide analysis [6–11]. Chromatographic methods have advantages such as high sensibility
17
18 77 and reliability, but the equipment for these methods is quite expensive, their productivity is
19
20 78 relatively low, and such methods require laborious and time-consuming preliminary sample
21
22 79 treatment. To make the testing of large number of samples cheaper and faster, chromatographic
23
24 80 analysis is accomplished by preliminary screening tests. The main purpose of preliminary
25
26 81 screening is to reduce the number of samples for confirmatory (chromatographic) analysis. This
27
28 82 wide screening is mainly focused on the most typical contaminants for the given territory, given
29
30 83 kind of samples, etc. For screening purposes, immunoassay methods are the most suitable
31
32 84 because of their specificity, sensitivity, rapidity, and low cost [12, 13]. Formerly, application of
33
34 85 enzyme-linked immunosorbent assay (ELISA) [14–16], strip-based immunoassays [17], and a
35
36 86 surface plasmon resonance [18] method have been reported for thiabendazole analysis, and
37
38 87 ELISA methods [19–21] have been applied for tetraconazole analysis. Besides,
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40 88 pseudoimmunoassay based on molecularly imprinted polymers has been reported for
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42 89 thiabendazole analysis [22]. No publications have been reported for fluorescence polarization
43
44 90 immunoassay (FPIA) of these compounds.
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51 91 The main advantages of FPIA as compared with other immunoassay techniques are its
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53 92 rapidity and simple manipulation (caused by one-stage homogeneous interaction of all analytical
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3 93 reactants and immediate changes of registered fluorescence polarization after immune binding).
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5 94 However, due to one-stage protocol without separation of formed immune complexes from
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8 95 initial reaction media the FPIA results are often sensitive to interfering matrix components, and
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10 96 so the assays in such cases should be accomplished by preliminary sample preparation
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12 97 procedures.

14 98 To date, preparation techniques for various kinds of samples have been successfully
15
16 99 adapted for determination of medicines, pesticides, mycotoxins, and other compounds using
17
18
19 100 FPIA [23, 24]. Recently, we developed FPIAs for triazophos and carbaryl analysis including
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21 101 sample preparation technique for wheat samples [25]. This research extends the frontiers of this
22
23 102 technique to other compounds and describes the first FPIAs methods for thiabendazole and
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25 103 tetraconazole.

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30 105 **Materials and methods**

32 106 **Reagents**

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35 107 Thiabendazole, tetraconazole, fluorescein isothiocyanate isomer I (FITC), 4-
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37 108 aminomethylfluorescein (AMF), aminofluorescein (AF), dicyclohexylcarbodiimide (DCC), N-
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39 109 hydroxysuccinimide (NHS), ethylenediamine dihydrochloride, 1,4-butylenediamine, and 1,6-
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41 110 hexamethylenediamine were purchased from Sigma Aldrich (St. Louis, MO, USA). Thin-layer
42
43 111 chromatographic (TLC) plates (silica gel) were purchased from Merck (Darmstadt, Germany).
44
45 112 All organic solvents and chemical reagents were of analytical reagent grade. Borate buffer (BB,
46
47 113 0.05 M, pH 8.6) with NaN_3 (0.01%) was used as a diluent for immunoreagents in FPIA.

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50 114 Haptens TN3C (3-[2-(1,3-thiazol-4-yl)-1*H*-benzimidazole-1-yl]propanoic acid), TN6C (3-
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52 115 [2-(1,3-thiazol-4-yl)-1*H*-benzimidazol-1-yl]hexanoic acid), and hapten DTPH (6-[2-(2,4-
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3 116 dichlorophenyl)-3-(1*H*-1,2,4-triazole-1-yl)propoxy]hexanoic acid) were earlier obtained by A.
4
5 117 Montoya (Universitat Politècnica de València, València, Spain). TN3C-BSA immunogen was
6
7 118 synthesized by the active ester technique [12] and used to obtain monoclonal antibodies against
8
9 119 thiabendazole (LIB-TN3C-13). DTPH-BSA was synthesized by the active ester technique [18]
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11 120 and used to obtain monoclonal antibodies against tetraconazole (LIB-DTPH-41).
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16 17 122 **Equipment**

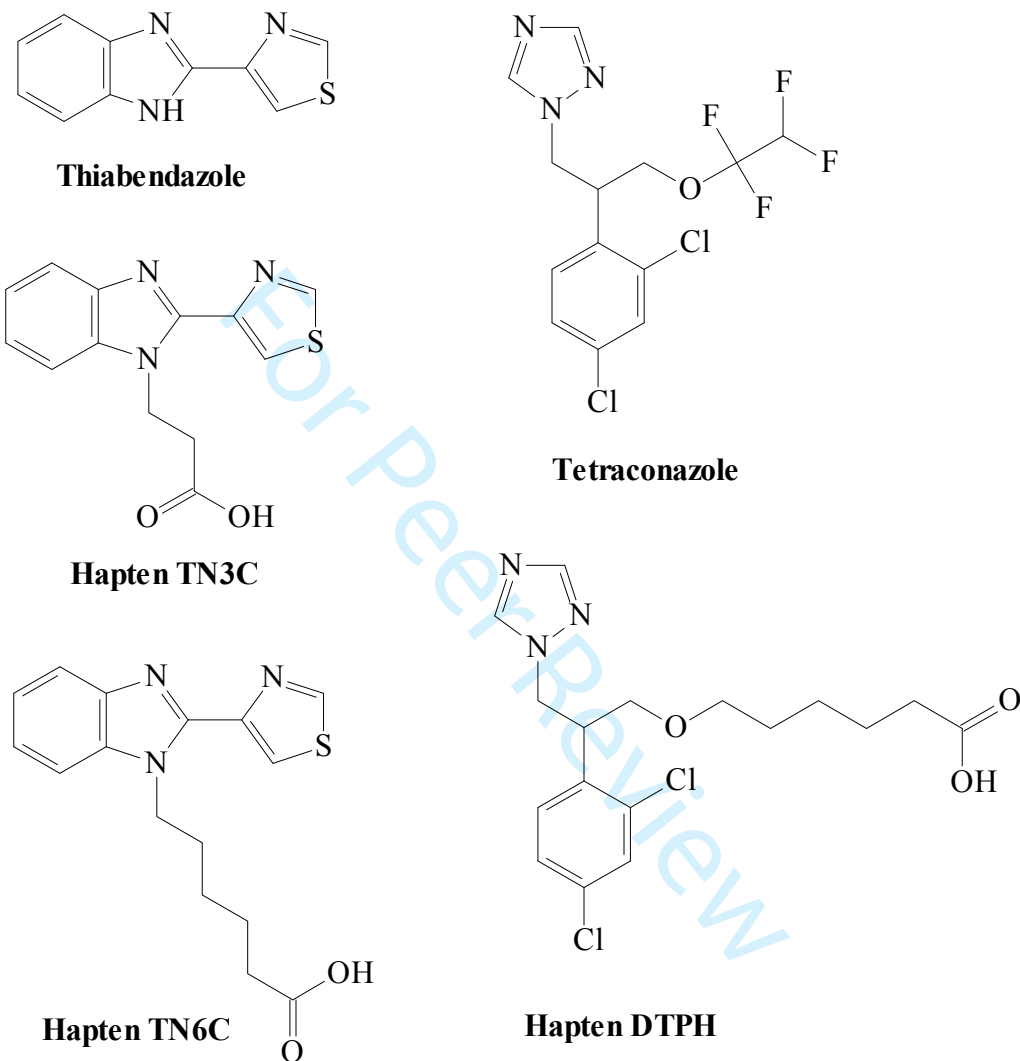
18
19 123 Measurements of fluorescence intensity and fluorescence polarization were performed
20
21 124 using a portable device, Sentry 200 (Ellie, Wauwatosa, WI USA). Data were processed using
22
23 125 Origin 8.5.1 software (OriginLab Corporation, Northampton, MA, USA). Mass-spectrometric
24
25 126 data were obtained using a tandem mass-spectrometer, Q-Exactive, coupled to a liquid
26
27 127 chromatograph DionexUltiMate 3000. Ionization of samples was performed using a HESI-II ion
28
29 128 source (Thermo Scientific, Waltham, MA, USA). Possible structures of fragment ions were
30
31 129 obtained using HighChem Mass Frontier 7.0 software from Thermo Scientific.
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36 37 131 **Synthesis of tracers**

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39 132 Labels purchased from Sigma Aldrich (AMF and AF) and labels synthesized by us
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41 133 [ethylenediaminefluoresceinthiocarbamyl (EDF), butylenediaminefluoresceinthiocarbamyl
42
43 134 ((CH₂)₄DF), and hexamethylenediaminefluoresceinthiocarbamyl ((CH₂)₆DF)] were used for the
44
45 135 synthesis of tracers. EDF was synthesized from FITC and ethylenediamine dihydrochloride as
46
47 136 described previously [26]. (CH₂)₄DF and (CH₂)₆DF were synthesized following the same
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49 137 technique used for EDF synthesis in which 1,4-butylenediamine and 1,6-hexamethylenediamine
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138 were used instead of ethylenediamine. Thiabendazole and tetraconazole do not contain functional
139 groups for synthesis of tracers, so their functionalized derivatives were used (Fig. 1).

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142

143 **Fig. 1** Chemical structures of analytes and haptens used for the preparation of tracers

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145 Tracers were synthesized as follows: 10 μmol of hapten were dissolved in
146 dimethylformamide (1 mL), then 4.3 mg of DCC (20 μmol) and 2.3 mg of NHS (20 μmol) were

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3 147 added to the solution. The reaction mixture was incubated for 12 h while stirring. The obtained
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5 148 precipitate was separated by centrifugation. Subsequently, 5 μmol of the fluorescent label were
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8 149 added to the supernatant, and the reaction mixture was mixed and incubated for 24 h.
9

10 150 Tracers were separated from the reaction mixtures by TLC. Tracers with diamine-FITC
11
12 151 labels were chromatographed with $\text{CHCl}_3:\text{CH}_3\text{OH}:\text{CH}_3\text{COOH}$ (80:16:1, v/v) as the mobile
13
14 152 phase. Bands of tracers with $R_f = 0.8$ were eluted from the TLC plate using methanol. Tracers
15
16
17 153 with the AF label were chromatographed with $\text{CHCl}_3:\text{CH}_3\text{OH}$ (5:1, v/v) and bands at $R_f = 0.4$
18
19 154 were eluted from the TLC plate. The separated fractions were additionally purified using TLC in
20
21 155 the mobile phase $\text{CHCl}_3:\text{CH}_3\text{OH}:\text{CH}_3\text{COOH}$ (80:16:1, v/v). After TLC separation, bands at $R_f =$
22
23 156 0.5 were eluted from the plates using methanol. For separation of the AMF-labeled tracers, the
24
25
26 157 mobile phase $\text{CHCl}_3:\text{CH}_3\text{OH}$ (8:1, v/v) was used; bands of tracers were at $R_f = 0.5$. The success
27
28 158 of syntheses and structures of tracers were confirmed by high-resolution tandem mass-
29
30 159 spectrometry coupled with high-performance liquid chromatography (HPLC-MS/MS).
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33 160

35 161 **FPIA procedure**

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37 162 The concentration of tracer solutions was estimated by fluorescence intensity (FI)
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39 163 measurement and its comparison with the FI for fluorescein. The FI of working solutions of
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41 164 tracers was 20 times higher than the FI of the buffer solution. The concentration of the working
42
43 165 solutions of the tracers was approximately 5 nM.
44
45

46
47 166 A series of dilutions were prepared to obtain antibody dilution curves. Each solution was
48
49 167 two times less concentrated than the previous solution. Aliquots (500 μL) of the diluted antibody
50
51 168 solution and 500 μL of the tracer working solution were mixed in each cuvette, and fluorescence
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3 169 polarization was measured. Dilution curves were produced using the results of the
4
5 170 measurements.

7
8 171 The choice of optimal dilution of antibodies and concentration of tracers for the most
9
10 172 sensitive PFIA was based on the presented experiments (Fig. 2, 3) in the accordance with
11
12 173 common practice of FPIA protocols development [23]. Other parameters of PFIA protocols such
13
14 174 as time of reactants incubation (2 min), pH of reaction mixture (8.5), nature and molarity of
15
16 175 buffer (0.05 M) were chosen on the basis of previous studies of fluorescein-based FPIA as
17
18 176 optimal for efficient immune interaction and fluorescence generation [27].

19
20
21 177 The FPIA procedure was performed as follows: 50 μL of a standard solution or a sample
22
23 178 were mixed in a cuvette with 500 μL of the tracer working solution, and fluorescence
24
25 179 polarization was measured. Results of the measurements of standard solutions were processed
26
27 180 using Origin software to obtain FPIA calibration curves. The time elapsed during the
28
29 181 measurement of the signal from a single sample was approximately 2–4 s.

30
31
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33 182 For experiments in selection of immunoreagents standard solutions were prepared in 10%
34
35 183 methanol, for experiments with wheat samples standard solutions were prepared in a mixture of
36
37 184 extractant and BB (1:7, v/v). Concentrations of thiabendazole standard solutions were 0.1, 1, 3.5,
38
39 185 10, 30, 100, 1000 ng/ml, concentrations of tetraconazole standard solutions were 1, 10, 35, 100,
40
41 186 300, 1000, 10000 ng/ml.

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45 187

46 47 188 **Data analysis**

48
49 189 The curves were plotted in coordinates "logarithm of concentration - mP" or "logarithm of
50
51 190 concentration - mP/mP_0 ", where mP is the measured fluorescence polarization, mP_0 is the
52
53 191 fluorescence polarization obtained for zero standard an optimized procedure of the analysis.

1
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3 192 These curves were approximated by a 4-parameter sigmoid equation:
4

5 193
$$Y = (A - D) / [1 + (x/C)^b] + D,$$

7 194 where A is the maximum value of the fluorescence polarization, D is the minimum value of the
8
9 195 fluorescence polarization, b is the slope of the curve at the IC₅₀ point, C (IC₅₀) is the analyte
10
11
12 196 concentration inhibiting the binding of antibodies to the tracer by 50%.

14 197 The limit of detection (LOD) was determined by performing the analysis of a blank
15
16 198 solution (solvent without analyte) 20 times. The LOD was calculated using a calibration curve as
17
18
19 199 the concentration corresponding to the difference between the average (blank) signal and three
20
21 200 times the standard deviation. 10% methanol was used as a blank solution to calculate LOD in standard
22
23 201 solutions and 8-fold diluted extract was used as a blank solution to calculate LOD in wheat samples.

25 202 The lower limit of quantification (IC₂₀) was calculated as the analyte concentration
26
27 203 inhibiting binding of the tracer with antibodies by 20%; the upper limit of quantification (IC₈₀)
28
29
30 204 was calculated as the analyte concentration inhibiting binding of the tracer with antibodies by
31
32 205 80%.

34 206 The cross-reactivity (CR) was calculated in accordance with Equation 1:

36 207
$$CR(\%) = (IC_{50}(\text{analyte}) / IC_{50}(\text{relative compound})) \times 100\% \quad (1)$$

38
39 208 where IC₅₀ is the concentration inhibiting binding by 50%.

41 209

43 210 **Obtaining of contaminated wheat grain**

46 211 The initial wheat grain preparations did not contain analytes, as shown by the HPLC-
47
48 212 MS/MS. The grain was ground in a homogenizer, then 1-gram samples of flour were
49
50
51 213 contaminated. Methanolic solutions of thiabendazole (1000 ng/ml) and tetraconazole (10,000
52
53 214 ng/ml) were used for this purpose. The obtained preparations contained 40, 100, 200, 300, 400
54
55 215 µg/kg of thiabendazole and 600, 1300, 1900, 2500 , 3200 µg/kg of tetraconazole. The

216 contaminated samples were left for 24 hours in a fume hood at a temperature of 22 ° C and a
217 humidity of 40%.

218

219 **Sample preparation**

220 Two milliliters of 70% methanol were added to 1 g of ground wheat. Samples were shaken
221 thoroughly and ultrasonicated for 30 min. The methanol fraction was diluted eight times with the
222 BB. The obtained precipitate was separated by centrifugation (5 min, 1400 g). The supernatant
223 was analyzed using FPIA and HPLC-MS/MS methods.

224

225 **HPLC-MS/MS-analysis**

226 A Hypersil Gold aQ column, Thermo Scientific, Waltham, MA, USA (150 × 2.1 mm i.d., 3
227 μm) with a Hypersil Gold aQ pre-column (10 × 2.1 mm i.d., 3 μm) was used for
228 chromatography. The column temperature was maintained at 30°C. The mobile phase consisted
229 of solvent A (0.1% formic acid in a mixture of water with acetonitrile 95:5, v/v) and solvent B
230 (0.1% formic acid in acetonitrile). The mobile phase gradient started at 5% B (0.0–2.0 min) and
231 increased to 95% B over 15.0 min, remained constant until 18.0 min, and was followed by
232 column equilibration to the initial conditions of 5% B (19.0–23.0 min). The flow rate of the
233 mobile phase was 0.5 mL/min, and the injection volume was 2 μL.

234 Mass spectrometric detection was performed under the following conditions: sheath gas
235 (nitrogen) flow rate, 0.4 L/min; auxiliary gas (nitrogen) flow rate, 0.1 L/min; sweep gas
236 (nitrogen) flow rate, 0.05 L/min; capillary voltage, 4.00 kV; capillary temperature, 270°C; and
237 auxiliary gas heater temperature, 280°C. The HESI-source was operated in the positive ion
238 mode. MS spectra were recorded under atmospheric pressure in the range of m/z 100–1500 Da,

239 the resolution was 35,000, and the isolation window was 5 ppm. MS/MS spectra were obtained
 240 using collision-induced dissociation. Collision energy for the tracers TN3C-EDF, TN6C-EDF,
 241 TN3C-AMF, TN6C-AMF, TN3C-AF, TN6C-AF, and DTPH-AF was 35%, and for the tracers
 242 DTPH-EDF, DTPH-(CH₂)₄DF, DTPH-(CH₂)₆DF, and DTPH-AMF was 20%. Collision energy
 243 for thiabendazole and tetraconazole in confirmation analysis of wheat samples was respectively
 244 40% and 35%. Conditions of quantitation and confirmation analysis are summarized in Table 1.

246 **Table 1** HPLC-MS/MS conditions for thiabendazole and tetraconazole analysis

Analyte	ESI mode	Precursor ion [M+H] ⁺ , Da	RT, min	Quantitation product ion (m/z)	Confirmation product ion (m/z)
Thiabendazole	positive	202.0439	3.9	175.0330	131.0608
Tetraconazole	positive	372.0294	10.1	158.9768	70.0405

248 Results and Discussion

249 FPIA development

250 The FPIA method is based on the competition between an antigen and a fluorescently-
 251 labeled antigen–tracer for a limited number of antibody binding sites. Immunoreagents should be
 252 selected so that the tracer can easily form the bond with the antibody and be displaced by the
 253 analyte. During the assay development, analytical characteristics of FPIA methods involving
 254 different tracers were compared. Structures of tracers were varied in two ways: by selecting
 255 different fluorescent labels for synthesis (EDF, AMF, AF) and by varying the fragment

256 connecting the antigen with the fluorescent label (the length of carbon bridge in TN3C and
 257 TN6C, the length of carbon bridge in EDF, (CH₂)₄DF, (CH₂)₆DF).

258 **Confirmation of the structures of tracers**

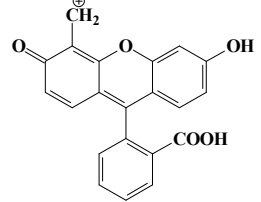
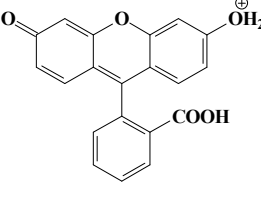
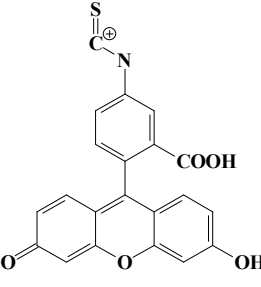
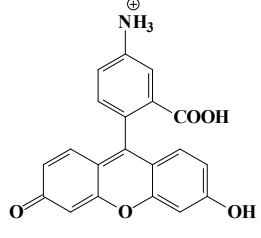
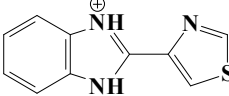
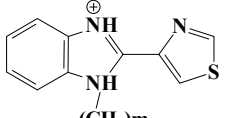
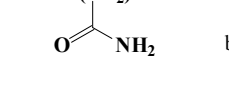
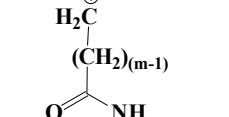
259 All the tracers used in this work were synthesized for the first time, so it was essential to
 260 confirm the success of syntheses and isolation of tracers from the reaction mixtures. HPLC
 261 coupled with tandem high-resolution mass spectrometry was used to identify the synthesized
 262 compounds.

263 Firstly, full scan mass spectra of tracers were obtained. Signals corresponding to singly and
 264 doubly protonated tracer molecules ($[M+H]^+$ and $[M+2H]^{2+}$) were present in positive ion mode
 265 spectra. Singly charged ions were used as precursor ions to obtain MS/MS spectra.

266 Peaks that were characteristic for fluorescent fragments of tracers were observed in
 267 MS/MS spectra of all the tracers. Peaks at m/z 345.0763 and 333.0763 were observed in spectra
 268 of tracers with the AMF label, peaks at m/z 390.0436 and 348.0872 were observed in spectra of
 269 tracers with diamine-FITC labels, and a peak at m/z 348.0872 was observed in spectra of tracers
 270 with the AF label. Molecular formulas and potential chemical structures are shown in Table 2.

271
 272 **Table 2** Accurate masses, molecular formulas and potential chemical structures of the product
 273 ions of tracers

Tracer molecule or tracer molecule section	m/z of product ion, Da	Molecular formulas of product ions	Potential structure of product ion
--	------------------------------	---------------------------------------	---------------------------------------

AMF-	345.0763	$C_{21}H_{13}O_5$	
	333.0763	$C_{20}H_{13}O_5$	
$(CH_2)_nDF^{-a}$	390.0436	$C_{21}H_{12}O_5NS$	
$(CH_2)_nDF^{-}, AF^{-}$	348.0872	$C_{20}H_{14}O_5N$	
TN3C-, TN6C-	202.0433	$C_{10}H_8N_3S$	
TN3C-AMF	273.0810	$C_{13}H_{13}N_4OS$	
TN6C-AMF	315.1280	$C_{16}H_{19}N_4OS$	
TN3C- EDF	115.0866	$C_5H_{11}N_2O$	

TN6C- EDF	157.1341	$C_8H_{17}N_2O$	
DTPH-EDF	428.1620	$C_{19}H_{28}Cl_2N_5O_2$	
DTPH-(CH ₂) ₄ DF	456.1933	$C_{21}H_{32}Cl_2N_5O_2$	
DTPH-(CH ₂) ₆ DF	484.2246	$C_{23}H_{36}Cl_2N_5O_2$	
DTPH-AMF	385.1198	$C_{17}H_{23}Cl_2N_4O_2$	
DTPH-AF	254.0252	$C_{11}H_{10}Cl_2N_3$	
^a n is the number of methylene groups; n=2, 4, 6 for tracers with EDF, (CH ₂) ₄ DF, and (CH ₂) ₆ DF labels, respectively;			
^b m is the number of methylene groups; n=2 and 5 for tracers with TN3C and TN6C haptens, respectively			

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275 In the MS/MS spectra of tracers synthesized from the haptens TN3C and TN6C, an intense
 276 peak of a product ion corresponding to the antigen fragment of the tracer molecules (m/z
 277 202.0433) was observed. Also, peaks corresponding to a carbon bridge between the antigen and
 278 the fluorophore of the tracer molecules (m/z 115.0866 and 157.1341) and peaks corresponding to
 279 a carbon bridge connected with the antigen section of the tracer (m/z 273.0810 and 315.1280)
 280 were obtained.

281 In the MS/MS spectra of the tracers synthesized from the DTPH hapten and the diamine-
 282 FITC labels, peaks corresponding to a carbon bridge connected with the antigen section of the

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3 283 molecule (m/z 428.1620, 456.1933, 484.2246) were observed. In the MS/MS spectrum of
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5 284 DTPH-AMF, a peak corresponding to the hapten (m/z 385.1198) was observed. In the MS/MS
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8 285 spectrum of DTPH-AF, a peak corresponding to the antigen section of the tracer molecule (m/z
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10 286 254.0252) was observed.

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13 14 288 **Selection of immunoreagents**

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17 289 **Tracers for thiabendazole analysis** Monoclonal antibodies obtained against the
18
19 290 immunogen TN3C-BSA by means of active ester method [14] were used to develop the FPIA of
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21 291 thiabendazole. TN3C is a thiabendazole derivative containing propanoic acid with a terminal
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23 292 carboxyl group as a spacer arm. TN3C and its homolog TN6C containing hexanoic acid with a
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25 293 terminal carboxyl group were used for the synthesis of tracers. AF, AMF, and EDF were used for
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27 294 the synthesis of tracers as fluorescent labels. Firstly, antibody dilution curves were obtained (Fig.
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29 300 2, A). Working concentrations of antibodies were chosen from the linear ranges of antibody
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31 295 dilution curves. Concentrations of antibodies were chosen for each tracer to compare them under
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33 296 the same conditions such that the difference between the maximum and minimum mP values on
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35 297 the calibration curve would be the same (70–80 mP). In subsequent experiments, antibodies were
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37 298 used in these concentrations. For illustrative purposes, data are shown at the coordinates of the
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39 300 plot of the mP/mP_0 versus analyte concentration, where mP is the measured fluorescence
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41 301 polarization, and mP_0 is the fluorescence polarization of the blank solution analyzed by the FPIA
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43 302 method (Fig. 2, B). Experiments were made in 3 or 4 replicates and errors varied from 1 to 5%.

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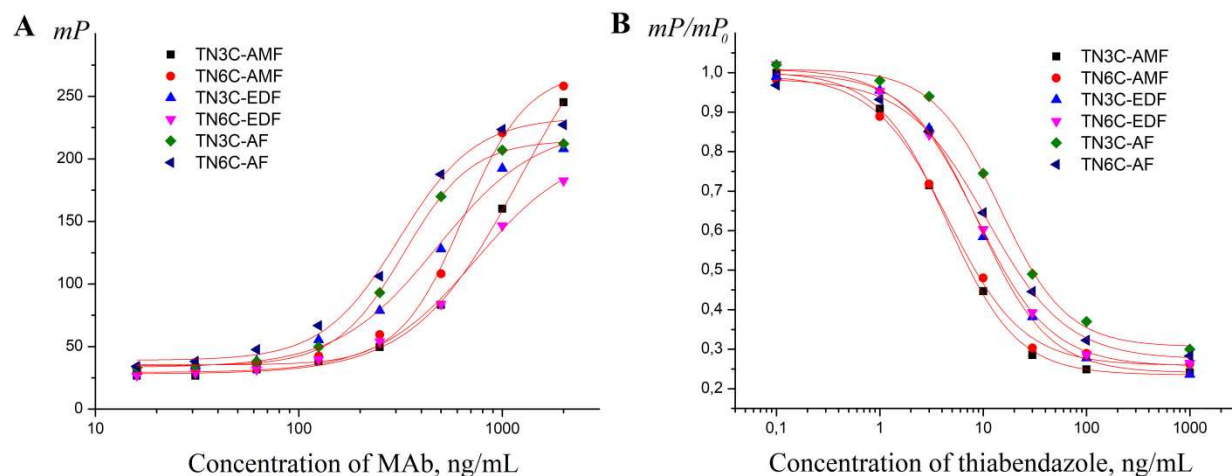


Fig. 2 (A) Antibody dilution curves and (B) calibration curves for thiabendazole determination using different tracers

When using tracers with the same fluorescent label and haptens having spacer arms of different lengths, the sensitivity of the FPIA method remained the same. Also, linear ranges of all the calibration curves accord to close concentrations; the minimal thiabendazole concentrations for these ranges with different tracers vary from 1.5 to 5.0 ng/mL, and the maximal thiabendazole concentrations – from 13 to 41 ng/mL (Table 3).

The choice of antibody concentrations for analysis was based on the need to ensure sufficient changes of analytic signal at the lowest antibody concentration and, by this way, to reach the best sensitivity. The standard deviations for the mP values in our experiment were from 0.8 to 2.5 mP. In comparison with the values of the analytical signal of 70-80 mP the deviations are less than 5%.

Using tracers with the AMF label yielded a slight advantage in sensitivity, whereas using TN6C-AMF reduced the amounts, and therefore the costs, of the antibodies required for the analysis. Consequently, TN6C-AMF was chosen for the subsequent experiments.

321

322 **Table 3** Characteristics of thiabendazole determination using different tracers

Tracer	Working concentration of antibodies, ng/mL	Linear range, ng/mL
TN3C-AMF	670	1.5–13
TN6C- AMF	500	1.5–15
TN3C-EDF	400	3.0–27
TN6C-EDF	670	2.8–27
TN3C-AF	280	5.0–41
TN6C-AF	250	3.2–38

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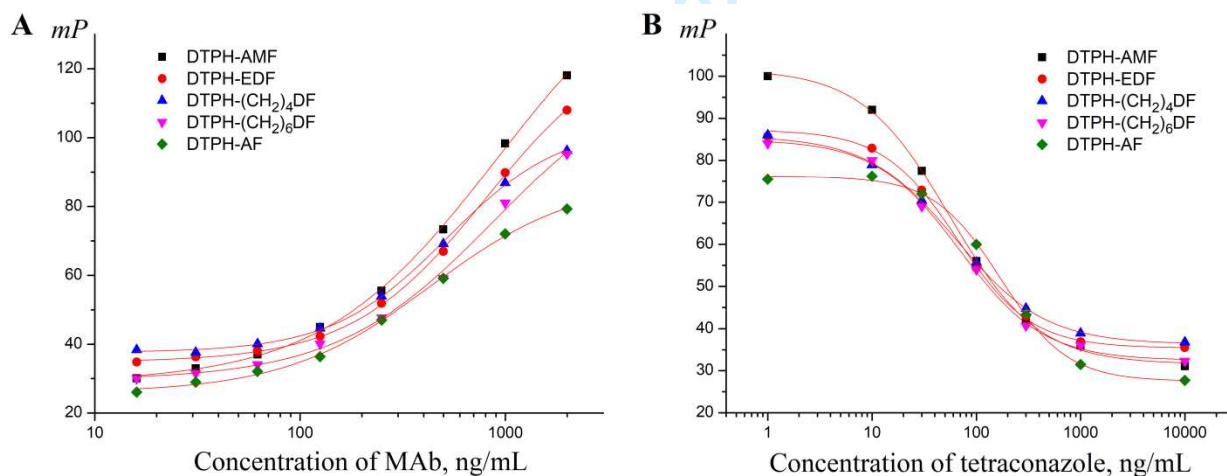
324 **Tracers for tetraconazole analysis**

325 Antibodies used in the development of the FPIA for tetraconazole were obtained against
 326 the synthesized by active ester method immunogen DTPH-BSA [20]. The hapten DTPH is a
 327 tetraconazole derivative containing a terminal carboxyl group. In this study, the hapten DTPH
 328 and the fluorescent labels AMF, EDF, (CH₂)₄DF, (CH₂)₆DF, and AF were used for the synthesis
 329 of tracers. Antibody dilution curves and calibration curves were obtained using the synthesized
 330 tracers (Fig. 3).

331 The difference between the maximum and minimum possible values of fluorescence
 332 polarization was calculated using the antibody dilution curves for each tracer. For DTPH-AMF,
 333 DTPH-EDF, DTPH-(CH₂)₄DF, DTPH-(CH₂)₆DF, and DTPH-AF this difference was 130, 100,
 334 70, 90, and 60 mP, respectively. The calibration curve for DTPH-AMF was obtained using a
 335 working solution of antibodies with a concentration of 1000 ng/mL to adjust the difference
 336 between the maximum and minimum mP values to be equal to 70–80 mP. Calibration curves

337 generated for other tracers were obtained using the same concentration of antibodies as used for
 338 DTPH-AMF. Other tracers bind antibodies significantly worse, and so the same range of
 339 fluorescence polarization values cannot be reached for them. Therefore, the conditions were
 340 standardized by selecting the same antibody concentration for all, namely 1000 ng/mL.

341 Calibration curves were prepared in the coordinates of plots of mP versus tetraconazole
 342 concentration for ease of comparison (Fig. 3, B). Experiments were made in 3 replicates and
 343 errors varied from 2 to 5%. The tracers DTPH-AMF, DTPH-EDF, DTPH-(CH₂)₄DF, and DTPH-
 344 (CH₂)₆DF yielded the same FPIA sensitivity (Table 4). However, using antibodies at a
 345 concentration of 1000 ng/mL with DTPH-AMF allowed operation in the range approximately
 346 from 30 to 100 mP ($\Delta mP = 70$), whereas using of any of the other tracers with antibodies in the
 347 same concentration reduced the ΔmP value to 40–50. Therefore, DTPH-AMF was chosen for
 348 FPIA development in wheat samples.



350
 351 **Fig. 3** (A) Antibody dilution curves and (B) calibration curves with different tracers for
 352 tetraconazole determination

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354 **Table 4** Characteristics of tetraconazole determination using different tracers

Tracer	Working concentration of antibodies, ng/mL	Linear range, ng/mL
DTPH-AMF	1000	16–210
DTPH-EDF	1000	21–180
DTPH-(CH ₂) ₄ DF	1000	20–250
DTPH-(CH ₂) ₆ DF	1000	20–250
DTPH-AF	1000	62–480

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356 **Characteristics of optimized FPIAs**

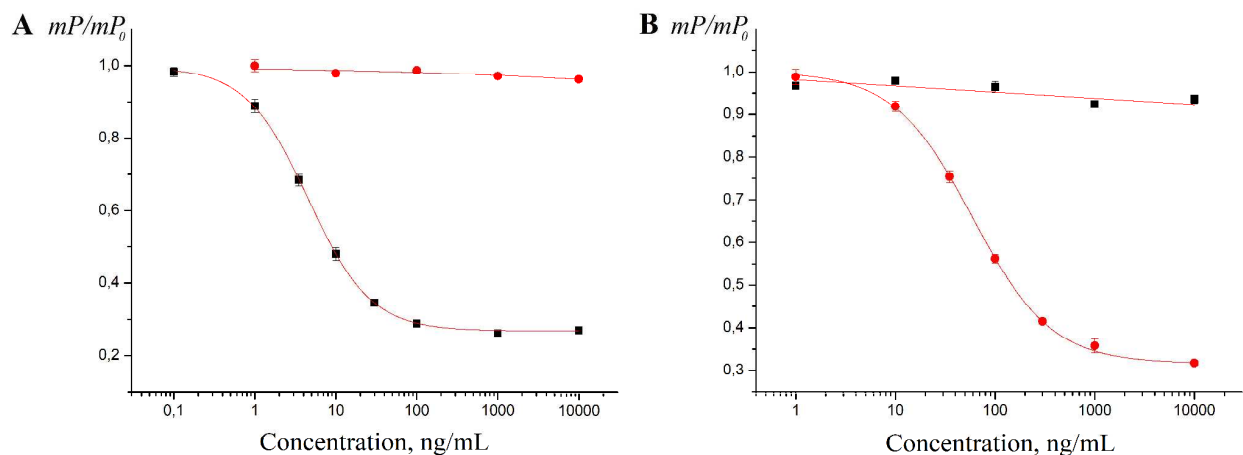
357 Calibration curves were prepared using the results of the analyses of thiabendazole and
358 tetraconazole standard solutions (Fig. 4). The linear range of determination for thiabendazole
359 was from 1.5 to 16 ng/mL, and the LOD was 1 ng/mL. The linear range of determination for
360 tetraconazole was from 16 to 210 ng/mL, and the LOD was 10 ng/ml. The specificity of the
361 developed methods was determined by comparison of the cross-reactivity with structurally
362 related compounds. Tetraconazole does not influence the interaction between anti-thiabendazole
363 antibodies and thiabendazole tracer (i.e. does not cause changes in the registered fluorescence
364 polarisation); thiabendazole does not influence the interaction between anti-tetraconazole
365 antibodies and tetraconazole tracer. The specificity of antibodies against thiabendazole was
366 investigated using the most similar compounds—benomyl and carbendazim. The cross-reactivity
367 for both compounds was less than 0.1%. The specificity of antibodies against tetraconazole LIB-
368 DTPH-41 was investigated using pesticides from the triazole class (Table 5). A high cross-
369 reactivity was observed for penconazole (35%) and cyproconazole (23%). Other tested triazoles
370 demonstrated negligible cross-reactivity. These results agree well with results obtained in

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3 371 previous reports [20] using the ELISA method. When normalized to a tetraconazole cross-
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5 372 reactivity of 100%, penconazole and cyproconazole demonstrated cross-reactivity at the levels of
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7 373 44% and 33%, respectively. Thus, the developed methods allowed a highly specific
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9 374 determination of thiabendazole and a less specific assay of tetraconazole because LIB-DTPH-41
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11 375 antibodies exhibited cross-reaction with other chemicals applied in agriculture.
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17 **Table 5** Cross-reactivity of anti-tetraconazole antibodies LIB-DTPH-41
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Compound	Cross-reactivity, %
Tetraconazole	100
Penconazole	35
Cyproconazole	23
Triadimefon	0.5
Propiconazole	0.3
Difenoconazole	<0.1
Tebuconazole	<0.1
Triadimenol	<0.1
Triticonazole	<0.1

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 380 **Fig. 4** (A) FPIA standard curve for thiabendazole (■) and cross-reactivity with tetraconazole (●);
 381 (B) FPIA standard curve for tetraconazole (●) and cross-reactivity with thiabendazole (■) (n=3)

382

383 Analysis of wheat samples

384 Thiabendazole and tetraconazole can occur in foodstuffs including cereals. The
 385 developed FPIA methods for thiabendazole and tetraconazole were applied for the analysis of
 386 wheat grain. The FPIA-adopted sample preparation technique for grain was published earlier
 387 [25].

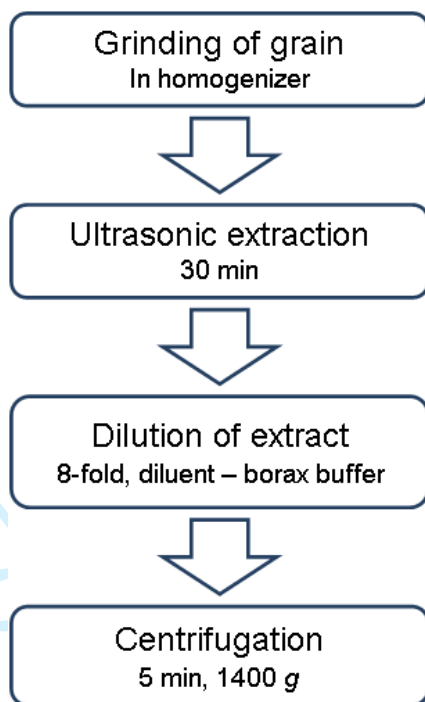
388 Grain is a complex matrix containing proteins, lipids, carbohydrates—starch,
 389 hemicelluloses, mucilages, and simple sugars—and mineral compounds. During sample
 390 preparation of grain, pesticides are usually extracted with organic solvents. The extracts obtained
 391 are then subjected to purification and concentration. Such sample preparation is not suitable for
 392 screening methods because it is time-consuming, whereas the main advantage of screening
 393 methods is their rapidity.

394 The sample preparation technique for FPIA in this research included ultrasonic extraction
 395 with 70% methanol, dilution of the extract with a buffer solution, and centrifugation. Scheme of
 396 sample preparation represented on Fig. 5. The choice of extractant was described in a recent

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3 397 manuscript [25] devoted to the FPIA of triazophos and carbaryl in wheat. Methanol was
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5 398 determined not to have a significant effect on the sensitivity of thiabendazole and tetraconazole
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10 400 The extract was diluted to eliminate matrix influence. The optimum extract dilution was
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12 401 determined in the following manner. An extract of uncontaminated wheat was prepared, and it
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14 402 was diluted 2-, 4-, 6-, and 8-fold. The diluted extracts were used to prepare standard solutions of
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16 403 analyte. The obtained standard solutions were analyzed via FPIA, and calibration curves were
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18 404 plotted using the results of this analysis. The calibration curves were compared with those
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20 405 obtained using the results of analysis of standard solutions in extractant diluted 2-, 4-, 6-, and 8-
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22 406 fold. Dilution of the extract was considered sufficient if the results analysis of standard solutions
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24 407 in diluted extracts and the corresponding diluted extractant nearly coincided, and the IC_{20} , IC_{50} ,
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26 408 IC_{80} , and mP_0 values differed insignificantly. The 8-fold dilution was concluded to be sufficient
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28 409 for the FPIA of thiabendazole and tetraconazole (Fig. 6, Table 6). Thiabendazole could be
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30 410 determined in wheat in the range from 40 to 500 $\mu\text{g}/\text{kg}$ with a LOD value of 20 $\mu\text{g}/\text{kg}$ using the
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32 411 developed FPIA method. Tetraconazole could be determined in the range from 600 to 3200
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34 412 $\mu\text{g}/\text{kg}$ with a LOD value of 200 $\mu\text{g}/\text{kg}$.

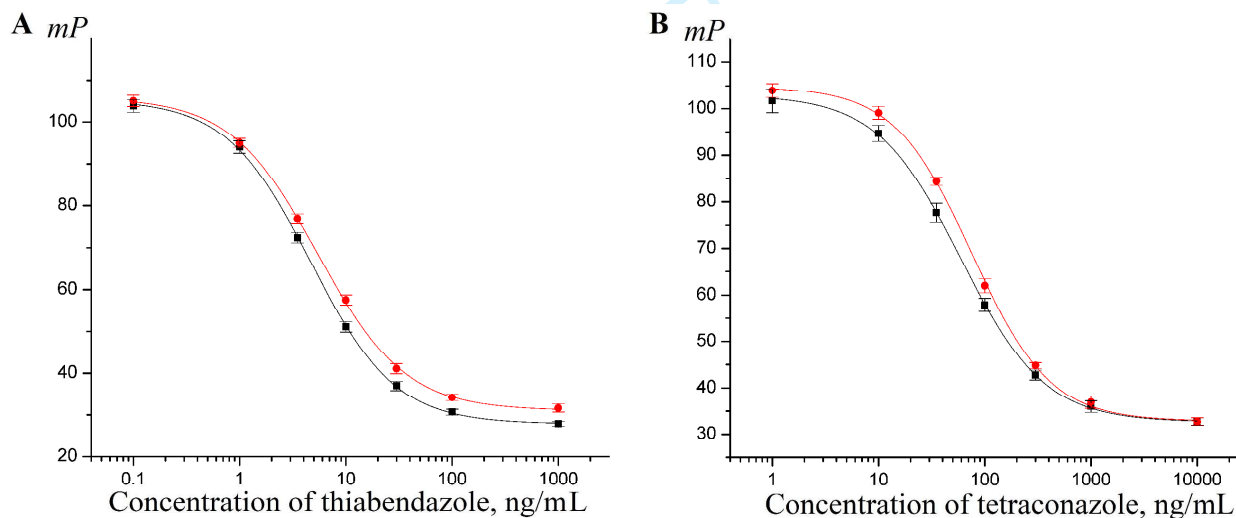
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414 **Fig. 5** Scheme of sample preparation

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417 **Fig. 6** FPIA standard curves for thiabendazole (A) and for tetraconazole (B) (■ - standards in

418 diluted extractant, ● – standards in diluted extract, n=3)

419 **Table 6** Parameters of sigmoidal fitting for calibration curves

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	(extractant)	(extract)	(extractant)	(extract)
IC20	1.3	1.5	17	24
IC50	4.7	5.4	59	76
IC80	17	20	210	245
R ²	0.999	0.999	0.999	0.999
ΔmP	77	75	70	72

420

421 Thus, the sample preparation technique developed earlier for the FPIA of triazophos and
422 carbaryl [25] was adapted for thiabendazole and tetraconazole. The dilution ratio of the extract
423 was determined for each compound. For triazophos, thiabendazole, and tetraconazole analysis
424 the minimum dilution ratio was 8, and for carbaryl analysis, it was 4. However, the sensitivity of
425 the carbaryl analysis was adequate for its determination below the existing MRLs even if an 8-
426 fold dilution was used, so the sample preparation technique for the FPIA of these four
427 compounds is identical.

428

429 **Recovery test**

430 Wheat grain used for the recovery test was not contaminated with pesticides. Wheat
431 samples were grinded; subsequently, they were spiked with thiabendazole and tetraconazole at
432 several concentrations, and the solvent was evaporated for 24 h. Preparation of spiked wheat
433 samples was conducted using the optimized conditions. The obtained diluted extracts were
434 analyzed in parallel by two methods—FPIA and HPLC-MS/MS.

435 The results of the recovery tests are presented in Table 6. The recoveries of thiabendazole
436 ranged from 71 to 86% by FPIA and from 83 to 89% by HPLC-MS/MS. The coefficient of

437 variation was less than 10% for the FPIA and less than 4% for the HPLC-MS/MS method (n=3).
 438 The recoveries of tetraconazole were from 60 to 77% by the FPIA and from 72 to 75% by the
 439 HPLC-MS/MS method; the coefficient of variation was less than 6% for the FPIA and less than
 440 2% for the HPLC-MS/MS method (n=3). The linear correlation between the results obtained by
 441 the two methods was observed. The regression equation for thiabendazole was $y = 0.819 x +$
 442 0.411 , with an R^2 value of 0.9985, and for tetraconazole it was $y = 0.989 x - 0.467$, with an R^2
 443 value of 0.9952. In general, the results obtained by the FPIA agreed with the results obtained by
 444 the HPLC-MS/MS method. Therefore, the developed methods of analysis using a fast and simple
 445 sample preparation technique were appropriate for the determination of thiabendazole and
 446 tetraconazole.

448 **Table 7** Analytical results and recoveries of thiabendazole and tetraconazole in wheat by the
 449 FPIA and HPLC-MS/MS methods.

Spiking level, μg/kg	FPIA		HPLC-MS/MS	
	Detected concentration ± SD, mg/kg	Recovery ± SD, %	Detected concentration ± SD, mg/kg	Recovery ± SD, %
Thiabendazole				
0	N. d. ^a	–	N. d.	–
40	34±3.4	86±8	36±1.1	89±3
100	75±6.8	75±7	88±0.7	88±1
200	144±4.5	72±2	171±6.8	86±3
300	213±6.8	71±2	248±2.8	83±1

400	272±4.6	68±1	340±7.9	85±2
Tetraconazole				
0	N. d.	–	N. d.	–
600	358±23	60±4	439±10	73±2
1300	900±31	69±2	936±9.1	72±1
1900	1470±57	77±3	1430±9.1	75±1
2500	1900±68	76±3	1850±12	74±1
3200	2450±68	77±2	2400±23	75±1
^a N. d. – not detected.				

450

451 **Conclusions**

452 FPIAs of thiabendazole and tetraconazole were developed for the first time. Tracers for FPIAs of
 453 these compounds were synthesized, and their structures were confirmed by HPLC-MS/MS. The
 454 influence of the structures of tracers on assay sensitivity was estimated. The sensitivity of the
 455 developed FPIAs depended on the structures of the fluorophores and antigen fragments of the
 456 tracer molecules and did not depend on the length of the bridge between them. FPIAs of
 457 thiabendazole and tetraconazole were applied for analysis of wheat using a sample preparation
 458 technique developed earlier that required less than an hour. This sample preparation technique
 459 has now been adapted for analysis of the four pesticides triazophos, carbaryl, thiabendazole, and
 460 tetraconazole. The LODs of thiabendazole and tetraconazole in wheat were 20 and 200 µg/kg,
 461 respectively. The linear range of thiabendazole determination was from 40 to 500 µg/kg, and the
 462 linear range of tetraconazole determination was from 600 to 3200 µg/kg. The results obtained by
 463 the proposed FPIA method exhibited a good correlation with the results obtained by the HPLC-

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3 464 MS/MS method. The developed methods are rapid, sensitive, and selective. Therefore, they are
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5 465 appropriate for high-throughput screening of thiabendazole and tetraconazole in wheat.
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10 467 **Acknowledgments**

11
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17 470 the obtained results.
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31 476 **Compliance with ethical standards**

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33 477 **Conflict of interest**

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35 478 The authors declare that they have no conflict of interest.
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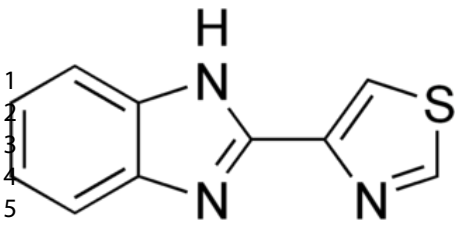
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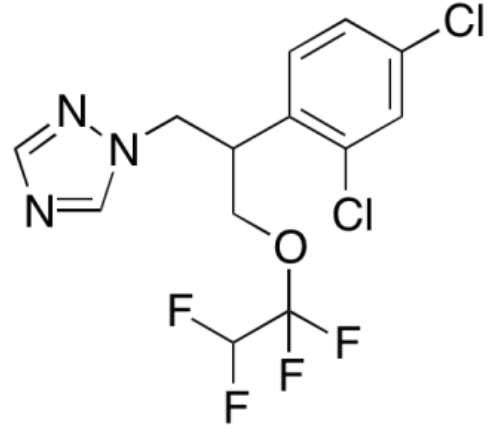
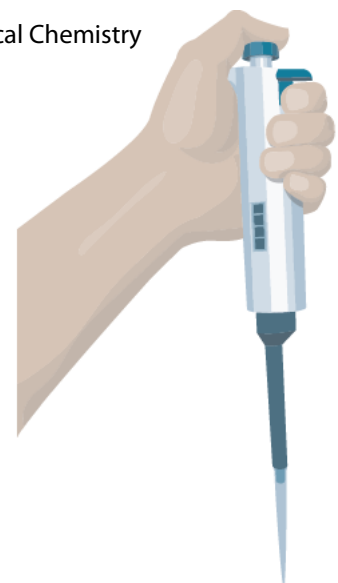
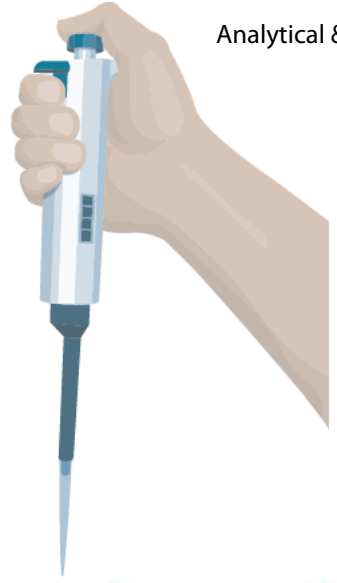
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For Peer Review



Thiabendazole



Tetraconazole



Two analytes – one procedure

- Unified sample preparation*
- Unified immune stage*
- Unified measurements of fluorescence polarization*

