

## TECHNICAL NOTE

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### Evaluation of Enzyme Immunoassay Performance Characteristics—Phencyclidine Example

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**ABSTRACT:** Four reagent formulations (three provided by a manufacturer; one prepared in-house by mixing equal volumes of two commercial reagents) are used for the assay of phencyclidine (PCP) in urine samples. Performance characteristics evaluated included assay precision and sensitivity at and near the assay cutoff concentration. Data resulting from the reagent prepared in-house are better than those using then commercially available formulations, and are comparable with those obtained using the recently available new commercial formulation.

**KEYWORDS:** toxicology, drug testing, enzyme immunoassay, phencyclidine

A drug of common abuse can often be tested by the same methodology using reagents provided by various manufacturers. Furthermore, a single manufacturer may supply various formulations suitable for different analyzer and test specificity requirements. Thus, it is interesting to compare the performance characteristics of various reagents.

Several studies on specific protocols have been reported [1–3]. We wish to report a comparative study using the Reply<sup>SM</sup> Automated Chemistry Analyzer (Olympus Corporation, Lake Success, NY) with three reagent formulations obtained from Syva Company (Palo Alto, CA) for the assay of phencyclidine (PCP) in urine samples. We also experimented a reagent formulation prepared in-house by 1:1 (v/v) mixing of two different Syva formulations.

Performance characteristics evaluated included assay precision and sensitivity at and near the assay cutoff concentration.

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## Materials and Methods

### Materials

The following assay kits were obtained from Syva Company (Palo Alto, CA): Emit<sup>®</sup> d.a.u.<sup>™</sup> Phencyclidine Assay kit, including Emit<sup>®</sup> d.a.u.<sup>™</sup> Phencyclidine Assay Antibody/Substrate Reagent A, Enzyme Reagent B, Emit<sup>®</sup> Drug Assay Buffer Concentrate, Emit<sup>®</sup> d.a.u.<sup>™</sup> Negative Calibrator, Emit<sup>®</sup> d.a.u.<sup>™</sup> Low Calibrator B, and Emit<sup>®</sup> d.a.u.<sup>™</sup> Medium Calibrator B; Emit<sup>®</sup> 700 Phencyclidine Assay kit, including Emit<sup>®</sup> 700 Phencyclidine Assay Antibody/Substrate Reagent 1, Enzyme Reagent 2, Emit<sup>®</sup> 700 Calibrator B, and Emit<sup>®</sup> 700 Control Set B (Positive Control B and Negative Control B); and Emit<sup>®</sup> II Phencyclidine Assay kit, including Emit<sup>®</sup> II Phencyclidine Assay Antibody/Substrate Reagent 1, Enzyme Reagent 2, Emit<sup>®</sup> Calibrator Level 0, and Emit<sup>®</sup> Calibrator A Level 1 (cutoff), Emit<sup>®</sup> Calibrator A Level 2 (high).

Reply<sup>™</sup> Automated Chemistry Analyzer from Olympus Corporation (Lake Success, NY) was used for all assays.

### Experimental Design

Four test protocols were conducted. Three of these protocols were those recommended by the manufacturer for their assay kits [4–6]: Emit<sup>®</sup> d.a.u.<sup>™</sup> Phencyclidine Assay (EMIT<sub>dau</sub>), Emit<sup>®</sup> 700 Phencyclidine Assay (EMIT<sub>700</sub>), Emit<sup>®</sup> II Phencyclidine Assay (EMIT<sub>II</sub>). The fourth protocol (EMIT<sub>mixed</sub>) used 1:1 (v/v) mixing of Antibody/Substrate Reagent A with Antibody/Substrate Reagent 1 and Enzyme Reagent B with Enzyme Reagent 2 from Emit<sup>®</sup> d.a.u.<sup>™</sup> Phencyclidine and Emit<sup>®</sup> 700 Phencyclidine Assay kits, respectively.

Sample, reagent, and diluent volumes recommended by the Reply<sup>™</sup> manufacturer were used for all protocols, specifically, the sample/diluent and the reagent/diluent (1-step; 2-step) volumes for the EMIT<sub>dau</sub>, EMIT<sub>700</sub>, and EMIT<sub>mixed</sub> protocols are: sample/diluent, 20/20; and reagent/diluent: 130/30, 130/30. The parallel parameters used for the EMIT<sub>II</sub> protocol are 8/5, 135/15, and 135/15, respectively.

## Results and Discussion

Perhaps the most important characteristic of a preliminary screen test is its ability to correctly differentiate samples containing the analyte at or above the cutoff concentration (25 ng/mL for PCP) from those containing less or no analyte. To meet this requirement, the assay should provide acceptable *sensitivity* and *precision* in the concentration range of concern. (Sensitivity is defined as the assay response per unit analyte concentration change, or simply, the slope of the dose-response plot.)

### Assay Sensitivity

Visual inspection of Fig. 1a (EMIT<sub>700</sub>) and 1b (EMIT<sub>dau</sub>) reveals the following performance characteristics:

1. The signal-dose plot of the EMIT<sub>700</sub> protocol in the 0 to 25 ng/mL range is linear with a steep slope, but becomes flat in the 25 to 50 ng/mL range; thus, the operation parameters recommended by Reply<sup>™</sup> manufacturer for the EMIT<sub>dau</sub> protocol cannot be used for the EMIT<sub>700</sub> assay;
2. The signal-dose plot of the EMIT<sub>dau</sub> is linear in the 0 to 50 ng/mL range with a response change (or “separation”) of only about 60 units, resulting in a smaller slope of the plot.

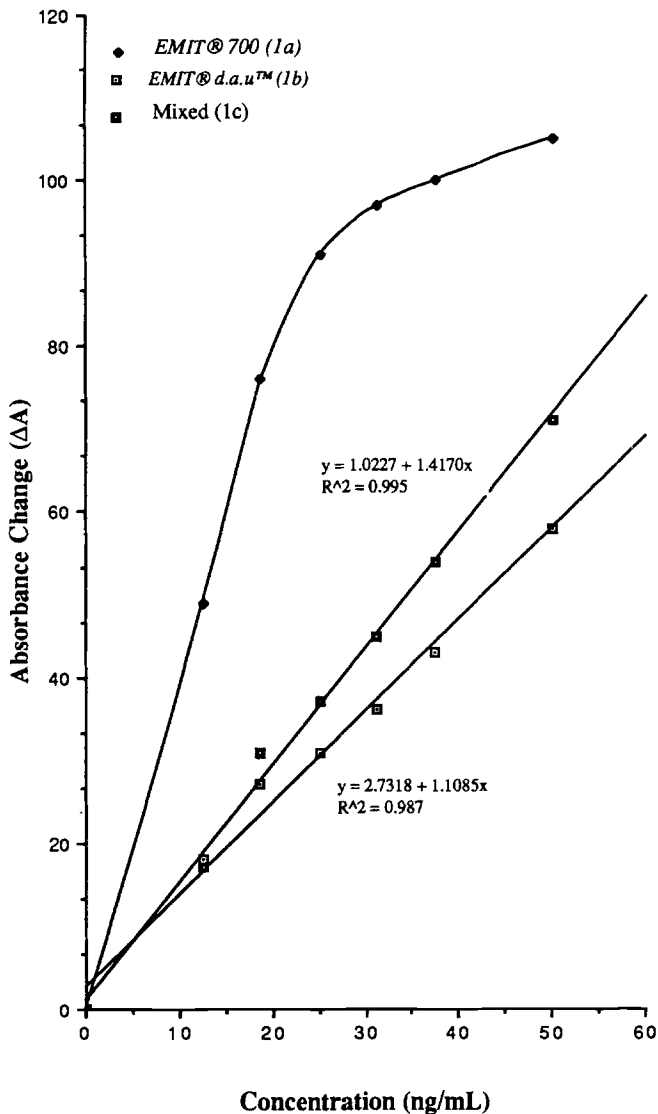


FIG. 1—Absorbance change vs. concentration plots using the *EMIT*<sub>700</sub> (1-a), the *EMIT*<sub>dau</sub> (1-b), and the *EMIT*<sub>mixed</sub> (1-c) protocols.

It was then rationalized that the analyte/reagent ratio of neither protocol is optimized for the targeted 0 to 50 ng/mL analyte concentration range, and that a more appropriate ratio may be obtained by mixing these two reagents. Indeed, the corresponding data (Fig. 1c) observed for the *EMIT*<sub>mixed</sub> protocol show improvement of assay sensitivity in increasing approximately 15 units of separation in the same concentration range.

#### Assay Precision

The sensitivity (of an assay protocol) that is required to effectively differentiate samples with a small concentration difference depends on the precision that can be achieved by

the assay; if the assay precision is high, the sensitivity become less critical. Parameters that can be effectively used for the evaluation of this aspect of performance characteristics are means and standard deviations (SD) obtained from samples with the analyte slightly above, at, or slightly below the cutoff concentration.

Individual data points obtained from within-run and between-day runs using the EMIT<sub>mixed</sub> protocols are compared in Tables 1 and 2. Individual data points obtained from within-run using the recently available product (Emit® II Phencyclidine Assay) are shown in Table 3, and compared with the corresponding data obtained from the EMIT<sub>mixed</sub> protocol in Fig. 2.

TABLE 1—*Within-run ΔA readings and statistical data (EMIT<sub>mixed</sub> protocol).*

|                      | Neg  | Concentration (ng/mL Phencyclidine) |      |      |      |      |      |
|----------------------|------|-------------------------------------|------|------|------|------|------|
|                      |      | 12.5                                | 18.8 | 25.0 | 31.3 | 37.5 | 50.0 |
| 1                    | 201  | 223                                 | 237  | 243  | 250  | 258  | 275  |
| 2                    | 206  | 224                                 | 235  | 243  | 251  | 260  | 276  |
| 3                    | 207  | 221                                 | 234  | 242  | 251  | 258  | 274  |
| 4                    | 205  | 225                                 | 236  | 245  | 253  | 261  | 277  |
| 5                    | 206  | 221                                 | 238  | 243  | 251  | 260  | 277  |
| 6                    | 204  | 224                                 | 237  | 241  | 252  | 261  | 277  |
| 7                    | 207  | 221                                 | 237  | 245  | 250  | 260  | 275  |
| 8                    | 206  | 222                                 | 237  | 243  | 250  | 259  | 277  |
| 9                    | 206  | 225                                 | 237  | 242  | 251  | 260  | 276  |
| 10                   | 203  | 221                                 | 236  | 242  | 249  | 259  | 279  |
| 11                   | 206  | 224                                 | 235  | 243  | 250  | 258  | 277  |
| 12                   | 206  | 223                                 | 238  | 243  | 250  | 259  | 275  |
| 13                   | 203  | 222                                 | 237  | 241  | 249  | 261  | 275  |
| 14                   | 204  | 222                                 | 237  | 242  | 251  | 260  | 277  |
| 15                   | 205  | 224                                 | 235  | 244  | 251  | 259  | 278  |
| Ave                  | 205  | 222                                 | 236  | 242  | 250  | 259  | 276  |
| SD                   | 1.69 | 1.47                                | 1.18 | 1.20 | 1.05 | 1.06 | 1.34 |
| %CV                  | 0.82 | 0.66                                | 0.50 | 0.49 | 0.42 | 0.40 | 0.48 |
| Difference from neg. | —    | 17                                  | 31   | 37   | 45   | 54   | 71   |

TABLE 2—*ΔA readings and statistical data of 12 batches run on 10 different days (EMIT<sub>mixed</sub> protocol).*

| Date                 | Neg  | Concentration (ng/mL Phencyclidine) |      |      |      |      |
|----------------------|------|-------------------------------------|------|------|------|------|
|                      |      | 12.5                                | 25.0 | 31.3 | 37.5 | 50.0 |
| 5/28                 | 180  | 203                                 | 213  | 221  | 235  | 248  |
| 5/28                 | 177  | 201                                 | 214  | 225  | 235  | 254  |
| 5/30                 | 181  | 203                                 | 216  | 223  | 235  | 248  |
| 5/31                 | 179  | 197                                 | 213  | 222  | 229  | 246  |
| 6/02                 | 177  | 192                                 | 206  | 216  | 223  | 240  |
| 6/13                 | 189  | 209                                 | 226  | 234  | 242  | 258  |
| 6/14                 | 187  | 205                                 | 221  | 231  | 238  | 255  |
| 6/16                 | 183  | 205                                 | 218  | 229  | 237  | 254  |
| 6/18                 | 186  | 205                                 | 219  | 229  | 239  | 257  |
| 6/19                 | 184  | 203                                 | 219  | 227  | 237  | 254  |
| 6/20                 | 184  | 203                                 | 216  | 228  | 241  | 253  |
| 6/20                 | 188  | 208                                 | 224  | 234  | 244  | 259  |
| Ave                  | 183  | 203                                 | 217  | 227  | 236  | 252  |
| SD                   | 4.14 | 4.60                                | 5.38 | 5.42 | 5.72 | 5.59 |
| %CV                  | 2.26 | 2.27                                | 2.48 | 2.39 | 2.42 | 2.21 |
| Difference from neg. | —    | 20                                  | 34   | 44   | 53   | 69   |

TABLE 3—Within-run  $\Delta A$  readings and statistical data (EMIT<sub>II</sub> protocol).

|                      | Neg  | Concentration (ng/mL Phencyclidine) |      |      |      |      |      |
|----------------------|------|-------------------------------------|------|------|------|------|------|
|                      |      | 12.5                                | 18.8 | 25.0 | 31.3 | 50.0 | 75.0 |
| 1                    | 163  | 175                                 | 186  | 196  | 205  | 232  | 247  |
| 2                    | 162  | 174                                 | 186  | 195  | 208  | 235  | 251  |
| 3                    | 161  | 176                                 | 185  | 193  | 207  | 235  | 250  |
| 4                    | 161  | 175                                 | 183  | 194  | 208  | 233  | 248  |
| 5                    | 161  | 173                                 | 184  | 194  | 208  | 233  | 251  |
| 6                    | 161  | 177                                 | 185  | 196  | 204  | 231  | 249  |
| 7                    | 160  | 177                                 | 183  | 193  | 206  | 232  | 248  |
| 8                    | 162  | 175                                 | 185  | 194  | 204  | 233  | 247  |
| 9                    | 161  | 176                                 | 182  | 192  | 206  | 233  | 246  |
| 10                   | 159  | 174                                 | 185  | 195  | 207  | 232  | 245  |
| 11                   | 161  | 175                                 | 183  | 193  | 203  | 232  | 250  |
| 12                   | 161  | 177                                 | 185  | 195  | 206  | 232  | 248  |
| 13                   | 161  | 173                                 | 184  | 193  | 205  | 230  | 246  |
| 14                   | 158  | 172                                 | 179  | 191  | 202  | 229  | 244  |
| Ave                  | 161  | 175                                 | 184  | 194  | 206  | 232  | 248  |
| SD                   | 1.23 | 1.59                                | 1.86 | 1.46 | 1.91 | 1.64 | 2.18 |
| %CV                  | 0.76 | 0.91                                | 1.01 | 0.75 | 0.93 | 0.71 | 0.88 |
| Difference from neg. | —    | 14                                  | 23   | 33   | 45   | 71   | 87   |

The mean and standard deviation data resulting from the EMIT<sub>dau</sub>, EMIT<sub>mixed</sub>, and the EMIT<sub>II</sub> protocols as shown in Table 4.

### Separation

The ability of an assay protocol to separate samples with different concentrations are evaluated on the overlapping (or nonoverlapping) characteristics between the following standards: 18.8 ng/mL (25% below cutoff) and 25 ng/mL (cutoff); and 25 ng/mL (cutoff) and 31.2 ng/mL (25% above cutoff). Corresponding data calculated from the EMIT<sub>dau</sub>, EMIT<sub>mixed</sub>, protocols and the EMIT<sub>II</sub> are shown in Table 5.

Data in Table 5 indicate that means of the paired standards are separated by more than 2 standard deviations with the EMIT<sub>mixed</sub> and EMIT<sub>II</sub> protocols, but not the EMIT<sub>dau</sub> protocol. Thus, the probability in differentiating (by the EMIT<sub>mixed</sub> and EMIT<sub>dau</sub> protocols) samples with concentrations 25% below or above the cutoff concentration from the cutoff is better than 97.72%. The EMIT<sub>mixed</sub> protocol was applied to test samples and found to correctly identify all positive samples as shown in Table 6.

### Deviation from Manufacturer's Recommended Procedure

The improved performance resulting from mixing two reagent formulations provided by the same manufacturer is itself an interesting observation worth reporting. From a practicing point of view, the established protocol was essential for effective use of the analyzer for PCP enzyme immunoassay at the time the study was conducted—EMIT<sub>II</sub> was then not yet available. Now that EMIT<sub>II</sub> formulation is available, this reported protocol provides an alternative approach with lower reagent cost.

For many laboratories, the most frequently encountered standards and criteria are those set forth by the National Laboratory Certification Program (NLCP) administered by the National Institute on Drug Abuse. On this matter, the NLCP has advised that

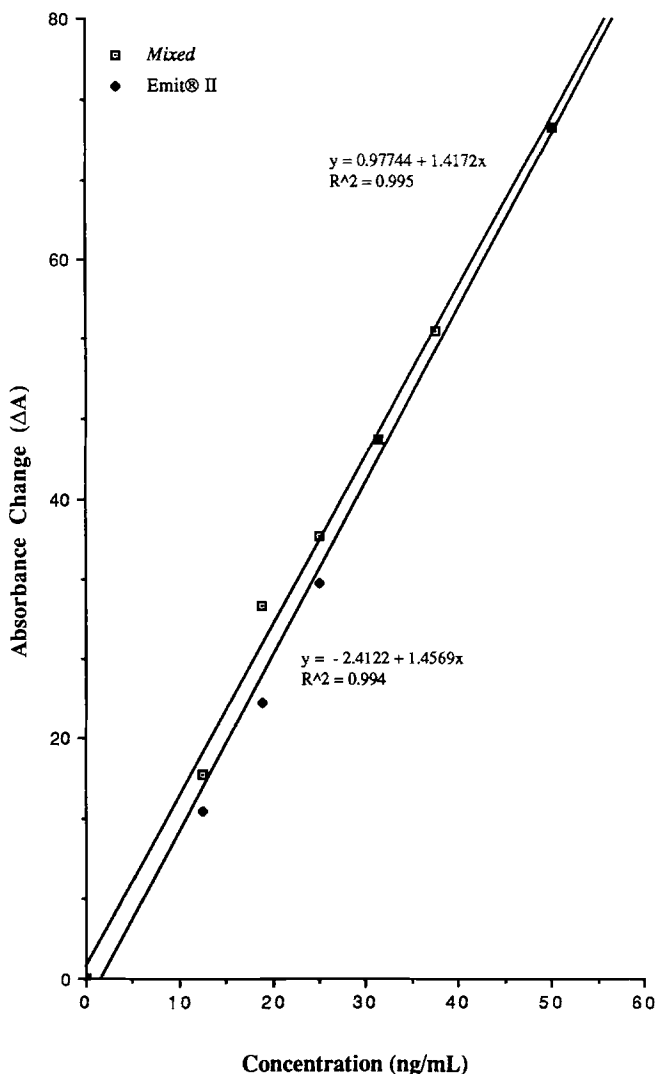


FIG. 2—Comparison of the  $EMIT_{mixed}$  and the  $EMIT_{II}$  protocols.

supporting changes to the manufacturers' procedures should, at a minimum, be characterized by data which defines the assay's linearity, precision, accuracy, detection limits, and specificity [8]. The laboratory should also demonstrate that the assay can differentiate between positive and negative specimens. Data reported herein were generated in the course of meeting these criteria.

**Conclusion**

With the operation parameters recommended by the Reply<sup>®</sup> manufacturer for the  $EMIT_{dau}$  protocol, the  $EMIT_{mixed}$  protocol provides better performance characteristics (linearity and sensitivity) than the  $EMIT_{dau}$  and  $EMIT_{700}$  protocols in the 0 to 50 ng/mL range. Samples with phencyclidine concentrations at 25% below or above the cutoff

TABLE 4—Mean and precision data of the  $EMIT_{duo}$ ,  $EMIT_{miscd}$ , and  $EMIT_{11}$  protocols.

| Protocol       | 0 ng/mL |      | 12.5 ng/mL |      | 18.8 ng/mL |      | 25 ng/mL |      | 31.3 ng/mL |      | 37.5 ng/mL |      | 50 ng/mL |      |
|----------------|---------|------|------------|------|------------|------|----------|------|------------|------|------------|------|----------|------|
|                | Ave     | SD   | Ave        | SD   | Ave        | SD   | Ave      | SD   | Ave        | SD   | Ave        | SD   | Ave      | SD   |
| $EMIT_{miscd}$ | 205     | 1.68 | 222        | 1.47 | 236        | 1.18 | 242      | 1.20 | 250        | 1.05 | 259        | 1.06 | 276      | 1.34 |
| $EMIT_{duo}$   | 194     | 3.05 | 213        | 2.66 | 221        | 1.96 | 224      | 2.20 | 230        | 1.31 | 237        | 1.51 | 252      | 1.52 |
| $EMIT_{11}$    | 161     | 1.23 | 175        | 1.59 | 184        | 1.86 | 194      | 1.46 | 206        | 1.91 | —          | —    | 232      | 1.64 |

<sup>a</sup>Means and standard deviations of 15 measurements.

<sup>b</sup>Means and standard deviations of 3 measurements for 0 and 50 ng/mL, and of 8 measurements for other concentrations.

<sup>c</sup>Means and standard deviations of 14 measurements.

TABLE 5—Comparison of critical separations obtained from the EMIT<sub>dau</sub>, the EMIT<sub>mixed</sub>, and the EMIT<sub>II</sub> protocols.

| Protocol              | Concentration and mean $\Delta A$ plus and/or minus 2 standard deviation (SD) |                           |                           |                             |
|-----------------------|---|---------------------------|---------------------------|-----------------------------|
|                       | 18.8 ng/mL<br>[Mean + 2 SD]   | 25 ng/mL<br>[Mean - 2 SD] | 25 ng/mL<br>[Mean + 2 SD] | 31.3 ng/mL<br>[Mean - 2 SD] |
| EMIT <sub>mixed</sub> | 238   | 240                       | 244                       | 248                         |
| EMIT <sub>dau</sub>   | 225   | 220                       | 228                       | 227                         |
| EMIT <sub>II</sub>    | 178   | 180                       | 188                       | 191                         |

TABLE 6—Application of the EMIT<sub>mixed</sub> protocol.

| Sample | GC/MS Conc. (ng/mL) | $\Delta A$ | EIA Result <sup>a,b</sup> |
|--------|---------------------|------------|---------------------------|
| 1      | 0                   | 202        | Negative                  |
| 2      | 0                   | 195        | Negative                  |
| 3      | 31                  | 246        | Positive                  |
| 4      | 0                   | 200        | Negative                  |
| 5      | 0                   | 198        | Negative                  |
| 6      | 32                  | 245        | Positive                  |
| 7      | 0                   | 205        | Negative                  |
| 8      | 0                   | 204        | Negative                  |
| 9      | 34                  | 245        | Positive                  |
| 10     | 0                   | 194        | Negative                  |
| 11     | 0                   | 205        | Negative                  |
| 12     | 32                  | 247        | Positive                  |
| 13     | 31.3 (Control)      | 249        | Positive                  |
| 14     | 37.5 (Control)      | 258        | Positive                  |
| 15     | 31.3 (Control)      | 251        | Positive                  |
| 16     | 37.5 (Control)      | 259        | Positive                  |
| 17     | 0                   | 204        | Negative                  |
| 18     | 0                   | 192        | Negative                  |
| 19     | 39                  | 256        | Positive                  |
| 20     | 37                  | 253        | Positive                  |

<sup>a</sup>Individual readings, mean, standard deviation, and %CV for the cutoff standards of the batch are: 237, 235, 231; 234; 3.06; and 1.31%, respectively. Thus, 234 was used as the cutoff value.

<sup>b</sup>Individual readings, mean, standard deviation, and %CV for the negative standards of the batch are: 204, 195, 199; 199; 4.51; and 2.27%, respectively.

standard (25 ng/mL) can be differentiated from the cutoff with a probability better than 97.72% (2 standard deviation). Results obtained using the EMIT<sub>mixed</sub> protocol are comparable with those obtained using the recently available EMIT<sub>II</sub> protocol.

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