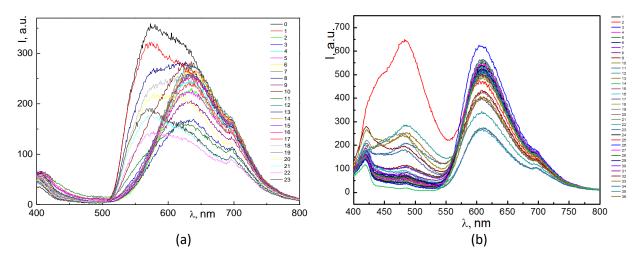
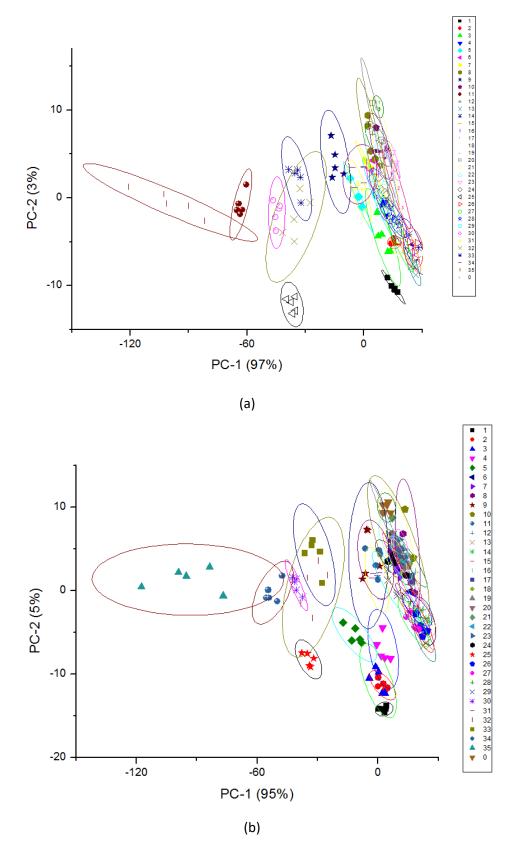
## Evaluation of discrimination performance in case for multiple non-discriminated samples: classification of honeys by fluorescent fingerprinting

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**Figure S1.** Fluorescence spectra of honey samples with added fluorophores: (**a**) intercalated thiazole orange (TO, ratio DNA-2 : TO = 1 : 105,  $\lambda_{ex}$  = 320 nm). With large amounts of the dye (TO:DNA ratios greater than 1:53), the intrinsic fluorescence of honey is quenched, and only a TO emission peak is observed in the spectrum. The largest number of discriminated groups was achieved with the TO:DNA-2 ratio of 1:53; (**b**) Ru(bpy)<sub>3</sub><sup>2+</sup> (50 µL of 1×10<sup>-5</sup> mol/L solution added to each well). The outlaying red spectrum belongs to a deeply colored sample.



**Figure S2.** Scores plots of the visible region reflection images of honey samples in the 96-well plates: (a) without added  $\text{Ru}(\text{bpy})_{3^{2^+}}$ , (b) in the presence of  $\text{Ru}(\text{bpy})_{3^{2^+}}$ . Conditions are the same as used in fluorescence studies.

Composition of the Blend (Ratio TO:DNA)	Number of Discriminated Groups of Points in the Scores Plot		
	by Full Spectrum (400-800 nm)	by One Peak	
		Shortwave	Longwave
Honey	7	8	_
Honey + DNA	11	10	_
Honey + DNA + TO (1:6)	10	8	5
Honey + DNA + TO (1:21)	8	5	5
Honey + DNA + TO (1:53)	9	_	12
Honey + DNA + TO (1:105)	4	_	6
Honey + DNA + TO (1:210)	4	_	5

**Table S1.** The number of discriminated groups of points in the scores plots of 23 honey samples with addedthiazole orange (TO) intercalated into DNA-2 (TO-DNA-2 fluorophore). The groups were consideredseparated if their confidence ellipses did not intersect

## Standard formulas for the calculation of confidence ellipses (based on the instruction to Origin software)

Assuming the pair of variables (X, Y) conforms to a bivariate normal distribution, we can examine the correlation between the two variables using a confidence ellipse. The confidence ellipse is centered at  $(\bar{x}, \bar{y})$  (for a given dataset  $(x_i, y_i)$ , i = 1, 2, ..., n, where x is the independent variable and y is the dependent variable), and the major semiaxis a and semiaxis b equal:

$$\bar{x} = \frac{1}{n} \sum_{i=1}^{n} x_i, \ \bar{y} = \frac{1}{n} \sum_{i=1}^{n} y_i,$$

$$a = c_{\sqrt{\frac{\sigma_x^2 + \sigma_y^2 + \sqrt{(\sigma_x^2 - \sigma_y^2) + 4r^2 \sigma_x^2 \sigma_y^2}}{2}}, \ b = c_{\sqrt{\frac{\sigma_x^2 + \sigma_y^2 - \sqrt{(\sigma_x^2 - \sigma_y^2) + 4r^2 \sigma_x^2 \sigma_y^2}}{2}},$$

where  $\sigma_i^2$  is the variance equal.

For a given confidence level of  $(1-\alpha)$  the confidence ellipse is defined as:

$$c = \sqrt{\frac{2(n+1)(n-1)}{n(n-2)}} \left( \alpha^{\frac{2}{2-n}} - 1 \right).$$

The inclination angle of the ellipse is defined as:

$$\beta = \frac{1}{2} \arctan \frac{2r \sqrt{\sigma_x^2 \sigma_y^2}}{\sigma_x^2 - \sigma_y^2}.$$

## Matlab function for relative position (RP) calculation (composed by the authors)

```
function[RP] = relative_position (sc1,sc2)
% sc1 - double [K1,2], where K1 is count of points in first class
% sc2 - double [K2,2], where K2 is count of points in second class
c1 = mean(sc1);% C1 - the center of the first group
c2 = mean(sc2);% C2 - the center of the second group
sc1 = get_border(sc1);% any border - ellipse or convex polygon
sc2 = get_border(sc2);
    % find A1,A2
     for i = 1:size(sc1,1)
         proj1(i) = (c1-c2)^{*}(sc1(i,:)-c2)'/(sqrt((c1-c2)^{*}(c1-c2)'));
     end
     for i = 1:size(sc2,1)
         proj2(i) = (c2-c1)^{*}(sc2(i,:)-c1)'/(sqrt((c1-c2)^{*}(c1-c2)'));
     end
    [~,a1] = min(proj1);
     a1 = sc1(a1,:);
    [~,a2] = min(proj2);
     a2 = sc2(a2,:);
     % find D1,D2
     di1 = find_d(a1,c1,c2);
```

```
di2 = find_d(a2,c1,c2);
% find RP
c1c2 = sqrt((c1-c2)*(c1-c2)');
c2d2 = sqrt((c2-di2)*(c2-di2)');
c1d1 = sqrt((c1-di1)*(c1-di1)');
RP = (c1c2-c2d2)/c1d1;
```

```
end
```

```
function[res] = find_d(a,ci1,ci2)
    eps = 10^{(-5)};
    n1 = ci2(1) - ci1(1);
    n2 = ci2(2) - ci1(2);
    if abs(n2) < eps \&\& abs(n1) < eps
         res = a;
     elseif abs(n2) < eps
         res = [a(1),ci1(2)];
     elseif abs(n1) < eps
         res = [ci1(1),a(2)];
     elseif abs(n2-1) < eps
         de1 = (a(1)*n1^2+a(2)*n1-ci1(2)*n1+ci1(1))/(n1^2+1);
         de2 = a(2)-n1^{*}(de1-a(1))/n2;
         res = [de1,de2];
     else
         e = -1/n1 - n1;
         g=n2-1/n2;
         f=ci1(2)/n2-ci1(1)/n1-n1*a(1)-n2*a(2);
         de1 = (a(1)*n1+a(2)*n2+n2*f/g)/(n1+n2*e/g);
         de2 = a(2)-n1^{*}(de1-a(1))/n2;
         res = [de1,de2];
     end
```

```
end
```

```
function[nodes] = get_border (X)%convex polygon
X = unique(X,'rows');
if size(X,1) == 2
nodes = X;
else
[~,i] = min(X(:,2));
nodes(1,:) = X(i,:);
X(i,:) = [];
for i = 1:size(X,1)
```

```
angle(i) = acot((X(i,2)-nodes(1,2))/(X(i,1)-nodes(1,1)));%acot == arc cotangent
        end
        [angle,i] = sort(angle,'descend');
        X = X(i,:);
        X(end+1,:) = nodes(1,:);
        nodes = [X(end-1,:);nodes;X(1,:)];
        i = 2;
        while i <= length(angle)+1
             if isLeft(nodes(end-2,:),nodes(end-1,:),nodes(end,:)) > 0
                 nodes(end+1,:) = X(i,:);
                 i = i + 1;
             else
                 nodes(end-1,:) = [];
             end
        end
    end
end
function[res] = isLeft(x0,x1,x2)
    res = ((x1(1)-x0(1))^*(x2(2)-x0(2)) - (x2(1)-x0(1))^*(x1(2)-x0(2)));
end
function[res] = dist(x)
    res = sqrt(x(1)^2+x(2)^2);
end
```