(11) **EP 1 529 849 A8** 

(12) CORRECTED EUROPEAN PATENT APPLICATION

Note: Bibliography reflects the latest situation

(15) Correction information:

Corrected version no 1 (W1 A2) INID code(s) 30

(48) Corrigendum issued on:

19.10.2005 Bulletin 2005/42

(43) Date of publication:

11.05.2005 Bulletin 2005/19

(21) Application number: 04447222.3

(22) Date of filing: 06.10.2004

(84) Designated Contracting States:

AT BE BG CH CY CZ DE DK EE ES FI FR GB GR HU IE IT LI LU MC NL PL PT RO SE SI SK TR Designated Extension States:

AL HR LT LV MK

(71) Applicant: De Ceuster NV 2860 sint-Katelijne-Waver (BE)

(72) Inventors:

 De Ceuster, Tom 2860 Sint-Katelijne-Waver (BE)

 Lievens, Bart 2860 Sint-Katelijne-Waver (BE) (51) Int CI.7: **C12Q 1/68** 

- Vanachter; Alfons 2860 Sint-Katelijne-Waver (BE)
- Krause, Matthew 2860 Sint-Katelijne-Waver (BE)
- Thomma, Bart Binnenhavens 5, 6709 PD Wageningen (NL)
- Cammue, Bruno
  Kasteelpark Arenberg 20, 3001 Heverlee (BE)
- (74) Representative: Brants, Johan P.E. et al De Clercq, Brants & Partners cv Edgard Gevaertdreef 10a 9830 Sint-Martens-Latem (BE)
- (54) Reliable method for assessing and reporting the risks of disease, contamination and losses caused by one or more microorganisms in a matrix or environment
- (57) The present invention relates to a method for assessing and reporting risks of disease, contamination or loss caused by one or more microorganisms in a matrix or an environment, said method comprising the steps of:
  - a) acquiring data regarding said one or more microorganisms and data regarding said matrix or said environment,
  - b) performing a risk analysis and assessment in order to determine a risk of disease, contamination or loss by said one or more microorganisms in said matrix or environment,
  - c) compiling the results of said risk analysis and outputting said results in a report, and

d) recommending suitable prevention, remediation or minimization strategies.

In a preferred embodiment, the invention comprises acquiring data regarding said microorganisms by detecting, identifying and quantifying one or more microorganisms or components thereof in said matrix or environment in a single assay, preferably using an array-based assay.