

## *Erratum*

# **Erratum to “Characterization of Selective Antibacterial Peptides by Polarity Index”**

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When I was using the Polarity-Index method a few days ago, I noticed differences in Table 3, as published.

However, it should be as in Table 3.

These changes do not affect neither the efficiency of the method nor any section of the paper.

I also noted the difficulty to compare information on the same APD2 database (November 2011), as it changes weekly. Therefore I am sending a file to support that test, in case the journal has backup for additional information. Otherwise I will keep this information for three years.

TABLE 3: Number of matches in a typical SCAAP sequence in each peptide database with single or multiple action on fungi, viruses, mammalian cells, Gram+/Gram- bacteria, cancer cells, insects, parasites, and sperms (see also Section 2.6) [7].

Total	Action	Fungi	Viruses	Mammalian cells	Bacteria	Cancer cells	Insects	Parasites	Sperms
879	Unique	0/77	0/22	0/10	62/743	1/16	0/2	0/9	0/0
2644	Multiple	72/638	6/122	33/205	152/1489	11/121	6/20	5/40	1/9