Human Keywords					
Strong effect		Strong neutral		Background	
Polymorphism	4,08%	Polymorphism	7,98%	Polymorphism	5,47%
Signal	3,47%	Phosphoprotein	6,55%	Phosphoprotein	3,41%
Membrane	2,91%	Cytoplasm	3,85%	Membrane	3,22%
Transmembrane	2,61%	Nucleus	3,19%	3D-structure	2,56%
Phosphoprotein	2,22%	Cytoskeleton	2,16%	Transmembrane	2,38%
Secreted	2,05%	Repeat	2,02%	Nucleus	2,33%
Acetylation	2,03%	Membrane	1,62%	Repeat	2,25%
3D-structure	1,94%	Transcription	1,08%	Cytoplasm	2,14%
Nucleus	1,80%	Metal-binding	0,85%	Glycoprotein	2,07%
Repeat	1,22%	Zinc	0,70%	Signal	1,67%

Tabelle1

Arabidopsis thaliana					
Strong effect		Strong neutral		Background	
Signal	5,30%	Membrane	3,19%	Membrane	3,95%
Membrane	3,97%	Transmembrane	2,60%	Transmembrane	3,37%
Transmembrane	3,75%	Nucleus	2,15%	Signal	2,26%
Secreted	2,98%	Signal	2,05%	Nucleus	2,02%
Antimicrobial	1,84%	Metal-binding	1,64%	Metal-binding	1,85%
Fungicide	1,84%	DNA-binding	1,46%	Transferase	1,85%
Metal-binding	0,92%	Repeat	1,32%	Repeat	1,81%
Nucleus	0,92%	Transcription	1,32%	Hydrolase	1,54%
Repeat	0,92%	Zinc	1,00%	Nucleotide-binding	1,42%
Transport	0,92%	Phosphoprotein	0,96%	ATP-binding	1,25%

Tabelle1

Escherichia coli					
Strong effect		Strong neutral		Background	
3D-structure	4,02%	3D-structure	4,20%	3D-structure	4,54%
Membrane	3,45%	Membrane	3,64%	Membrane	3,96%
Transmembrane	2,87%	Signal	3,08%	Transmembrane	3,17%
Ribonucleoprotein	1,72%	Transmembrane	3,08%	Transport	2,45%
Signal	1,72%	Repeat	2,24%	Cytoplasm	2,14%
Cytoplasm	1,15%	Transport	2,24%	Metal-binding	2,06%
DNA-binding	1,15%	Periplasm	1,40%	Transferase	1,90%
Toxin	0,57%	Cytoplasm	1,12%	Signal	1,63%
Transcription	0,57%	Hydrolase	1,12%	Nucleotide-binding	1,62%
tRNA-binding	0,57%	Nucleotide-binding	1,12%	Hydrolase	1,57%