							GO:0015979   photosynthesis
							GO:0018298   protein-chromophore linkage
							GO:0009644   response to high light intensity
							GO:0055114   oxidation_reduction process
							GO:0009414   response to water deprivation
							GO:1901700   response to oxygen–containing compound
							GO:0098542   defense response to other organism
							GO:0098869   cellular oxidant detoxification
							GO:0042744   hydrogen peroxide catabolic process
							GO:0006979   response to oxidative stress
							GO:0016036   cellular response to phosphate starvation
							GO:0010035   response to inorganic substance
							GO:0010200   response to chitin
							CO:0006469   protoin phosphorylation
							GO:0009247   glycolipid biosynthetic process   GO:0009247   glycolipid biosynthetic process   10
							GO:0009723   response to ethylene
							GO:0016311   dephosphorylation
							GO:0015608 Liporgapio anion transport
							GO:0019951   negative regulation of endopeptidas of the control of
							GO:0006720   isoprenoid metabolic process
							GO:0008299   isoprenoid biosynthetic process
							GO:0003333   amino acid transmembrane transport
							GO:0006633   fatty acid biosynthetic process
							GO:0009834   plant-type secondary cell wall biogenesis
							GO:0033692   cellular polysaccharide biosynthetic process
							GO:0046274   lignin catabolic process
							GO:0045491   xylan metabolic process
							GO:0071555   cell wall organization
							GO:0007010   cytoskeleton organization
							GO:0007018   microtubule-based movement
<u> </u>	212		47.	<u>.</u> 5	9.10		J ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '
Cluster1	Cluster2	Cluster3	Cluster4	Cluster5	Cluster6	Cluster7	
Ō	Ö	Ö	Ö	Ō	Ō	Ö	