Annotation in Gene Ontology database

Categories	%
Biological Process	7.47
Cellular Componen	8.76
Molecular Function	7.89
no hit	75.89

Biological Process	%
metabolic process	35.04
cellular process	12.53
biological regulation	11.56
response to stimulus	8.88
localization	6.67
developmental process	3.03
cellular component organication or biogenesis	2.29
multicellular organismal process	1.15
other	2.35
unclassified	16.49

Cellular Component	%
cell part	46.92
organelle	23.78
membrane part	8.16
ptotein-containing complex	6.18
membrane	5.67
organelle part	2.8
extracellular region	2.42
cell junction	1.01
other	0.49
unclassified	2.57

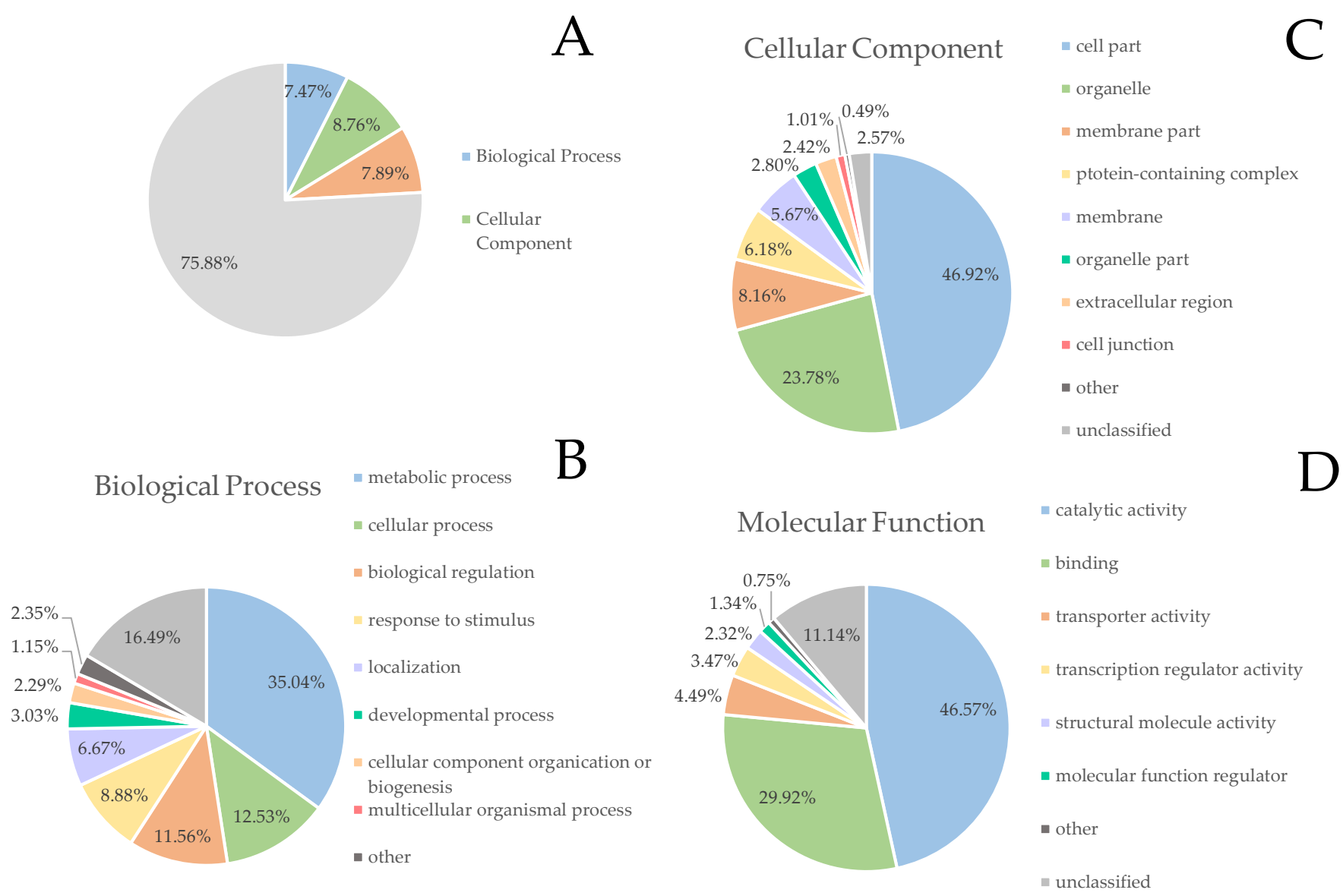| Molecular Function	%
catalytic activity	46.57
binding	29.92
transporter activity	4.49
transcription regulator activity	3.47
structural molecule activity	2.32
molecular function regulator	1.34
other	0.75
unclassified	11.14

Figure S1. Gene Ontology (GO) annotation of the assembled two lisianthus cultivars. All GO terms are grouped into three ontology (**A**) List of the GO terms belonging to biologilal process (**B**) List of the GO terms belonging to cellular component (**C**) List of the GO terms belonging to molecular function (**D**)