GO annotation for DSD-cooh proteins

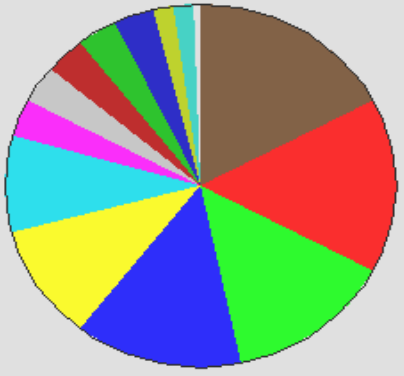
Functional Categorization by Annotations for : GO Cellular Component

- cellular component unknown: 20.5% (raw value = 9)
- other cellular components: 13.6% (raw value = 6)
- nucleus: 11.4% (raw value = 5)
- other membranes: 11.4% (raw value = 5)
- other intracellular components: 11.4% (raw value = 5)
- chloroplast: 9.1% (raw value = 4)
- other cytoplasmic components: 9.1% (raw value = 4)
- mitochondria: 6.8% (raw value = 3)
- ribosome: 2.3% (raw value = 1)
- cell wall: 2.3% (raw value = 1)
- extracellular: 2.3% (raw value = 1)

Functional Categorization by Annotations for : GO Molecular Function

- molecular function unknown: 23.8% (raw value = 10)
- DNA or RNA binding: 14.3% (raw value = 6)
- other molecular functions: 14.3% (raw value = 6)
- protein binding: 9.5% (raw value = 4)
- nucleic acid binding: 9.5% (raw value = 4)
- transporter activity: 9.5% (raw value = 4)
- other binding: 7.1% (raw value = 3)
- structural molecule activity: 4.8% (raw value = 2)
- other enzyme activity: 2.4% (raw value = 1)
- receptor binding or activity: 2.4% (raw value = 1)
- transcription factor activity: 2.4% (raw value = 1)

Functional Categorization by Annotations for : GO Biological Process



- biological process unknown: 17.2% (raw value = 10)
- other metabolic processes: 15.5% (raw value = 9)
- other cellular processes: 13.8% (raw value = 8)
- other physiological processes: 13.8% (raw value = 8)
- protein metabolism: 10.3% (raw value = 6)
- cell organization and biogenesis: 8.6% (raw value = 5)
- transport: 3.4% (raw value = 2)
- transcription: 3.4% (raw value = 2)
- developmental processes: 3.4% (raw value = 2)
- other biological processes: 3.4% (raw value = 2)
- electron transport or energy pathways: 3.4% (raw value = 2)
- response to stress: 1.7% (raw value = 1)
- response to abiotic or biotic stimulus: 1.7% (raw value = 1)

http://www.arabidopsis.org/servlet/Search