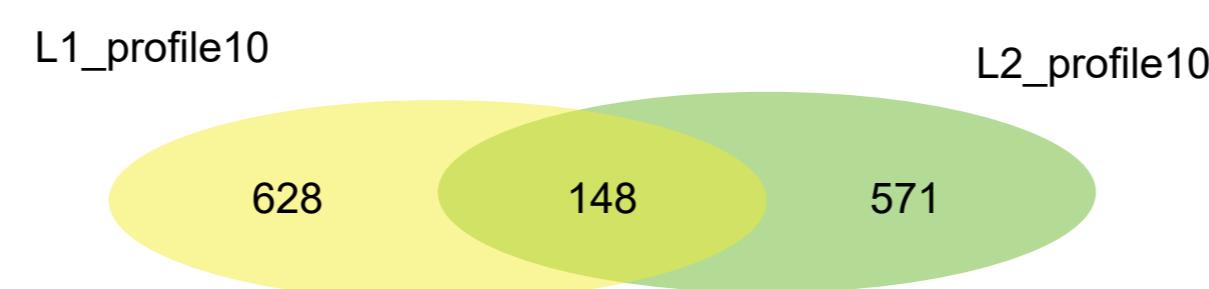
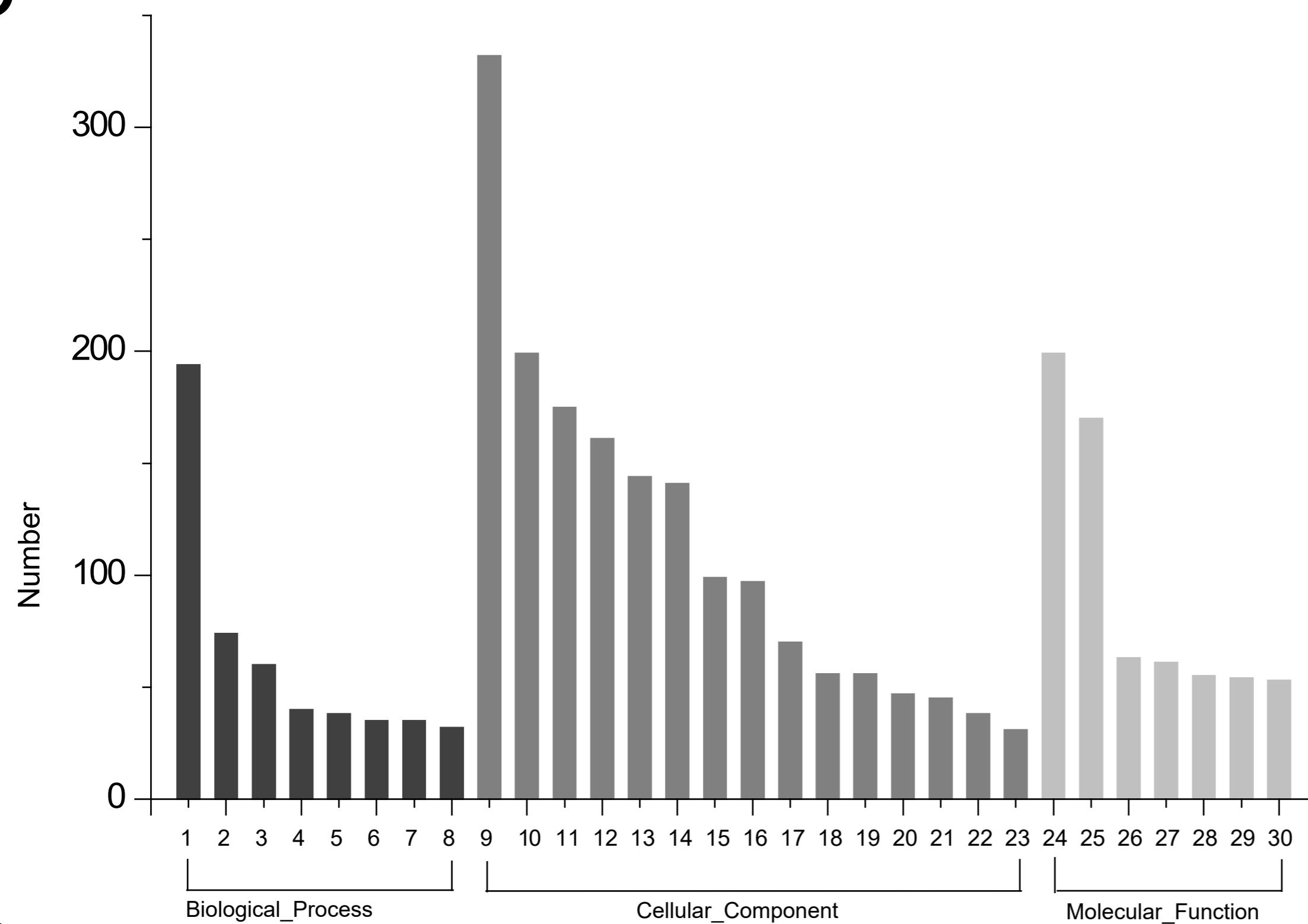
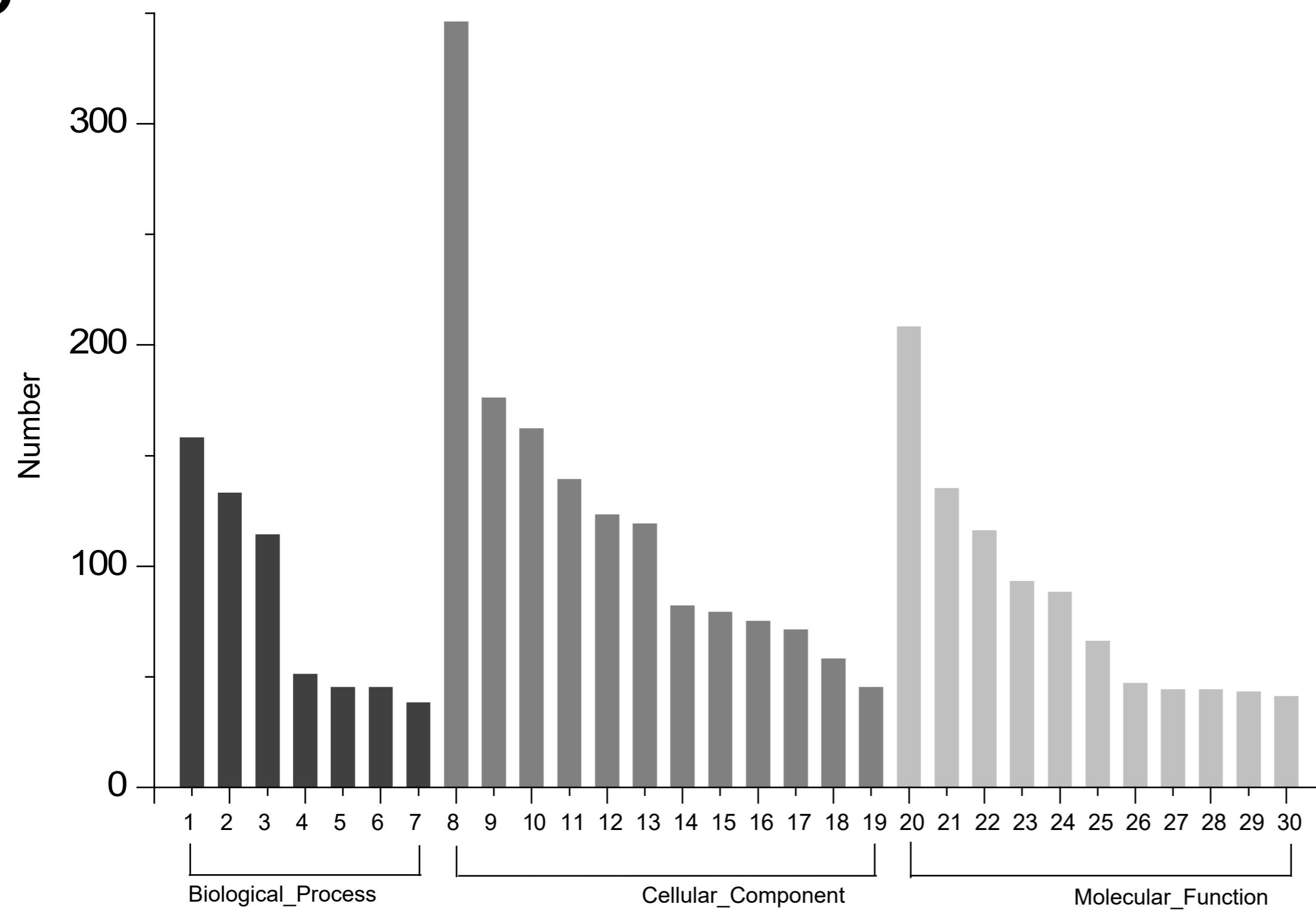


a**b**

- 1:biological_process
- 2:regulation of transcription, DNA-templated
- 3:transcription, DNA-templated
- 4:oxidation-reduction process
- 5:response to water deprivation
- 6:response to abscisic acid
- 7:response to salt stress
- 8:defense response to bacterium
- 9:nucleus
- 10:cytoplasm
- 11:integral component of membrane
- 12:plasma membrane
- 13:chloroplast
- 14:cytosol
- 15:membrane
- 16:mitochondrion
- 17:extracellular region
- 18:plasmodesma
- 19:cellular_component
- 20:vacuole
- 21:Golgi apparatus
- 22:endoplasmic reticulum
- 23:chloroplast stroma
- 24:molecular_function
- 25:protein binding
- 26:zinc ion binding
- 27:DNA binding
- 28:ATP binding
- 29:transcription factor activity, sequence-specific DNA binding
- 30:metal ion binding

c

- 1:biological_process
- 2:regulation of transcription, DNA-templated
- 3:transcription, DNA-templated
- 4:response to abscisic acid
- 5:response to salt stress
- 6:protein phosphorylation
- 7:negative regulation of transcription, DNA-templated
- 8:nucleus
- 9:cytoplasm
- 10:plasma membrane
- 11:chloroplast
- 12:integral component of membrane
- 13:cytosol
- 14:membrane
- 15:cellular_component
- 16:plasmodesma
- 17:mitochondrion
- 18:extracellular region
- 19:vacuole
- 20:protein binding
- 21:molecular_function
- 22:DNA binding
- 23:transcription factor activity, sequence-specific DNA binding
- 24:ATP binding
- 25:metal ion binding
- 26:zinc ion binding
- 27:sequence-specific DNA binding
- 28:protein serine/threonine kinase activity
- 29:kinase activity
- 30:transcription regulatory region DNA binding