
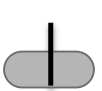
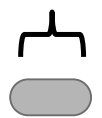





<b>SNP</b>		Identifier: RS number, example “rs1234”. Used to refer to a known and documented SNP whose position can be retrieved from the knowledge database.
<b>Position</b>		Identifier: Chromosome and basepair location, example “chr1:234”. Used to refer to any single genomic location, such as a single nucleotide polymorphism (SNP), single nucleotide variation (SNV), rare variant, or any other position of interest.
<b>Region</b>		Identifier: Chromosome and basepair range, example “chr1:234-567”. Used to refer to any genomic region, such as a copy number variation (CNV), insertion/deletion (indel), gene coding region, evolutionarily conserved region (ECR), functional region, regulatory region, or any other region of interest.
<b>Gene</b>		Identifier: Name, example “A1BG” or “ENSG00000121410”. Used to refer to a known and documented gene, whose genomic region and associations with any pathways, interactions or other groups can be retrieved from the knowledge database.
<b>Group</b>		Identifier: Name, example i.e. “lipid metabolic process” or “GO:0006629”. Used to refer to a known and documented pathway, ontological group, protein interaction, protein family, or any other grouping of genes, proteins or genomic regions that was provided by one of the external data sources.
<b>Source</b>		Identifier: Name, example “GO”. Used to refer to a specific external data source.