Basic Analysis

- Amino Acid Translation Element
- Annotate with UQL Element
- CD-Search Element
- Collocation Search Element
- Export PHRED Qualities Element
- Fetch Sequences by ID From Annotation Element
- Filter Annotation by Name Element
- Filter Annotations by Qualifier
- Find Correct Primer Pairs Element
- Find Pattern Element
- Find Repeats Element
- Gene-by-gene approach report
- Get Sequences by Annotations Element
- Group Primer Pairs Element
- Import PHRED Qualities Element
- Intersect Annotations Element
- Local BLAST Search Element
- Merge Annotations Element
- ORF Marker Element
- Remote BLAST Element
- Sequence Quality Trimmer Element
- Smith-Waterman Search Element