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retTaxonRvNameResponse	
getTaxonByNameResponse getESTsThatAreInAGivenESTLibraryReturn	
GetPublicationsBvSearchResponse	į
GetPublicationsBySearchResponse getEstGenomeMatchReturn	
getRelaTypesReturn	
doScoobyByIdResponse	
whatComponentExpressGeneReturn	
getNeuroBlast1Response	
getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameMCLClustersAsProteinsWithAGivenGOAnnotationReturn	<u> </u>
RNAsuboptSOAPResponse	<u> </u>
getNumberOfResultsResponse	
getMaxBatchSizeReturn searchCitationsResponse	
BlastofasResponse	
SimpleSearchResponse	
net Organisms Response	
getÖrganismsResponse getAllELMsResponse	
soap_create_modelResponse	
RetrievePairwiseScoresResponse	<u>;</u>
MOBYSHoundLocusIDFromGiReturn	i i

name	count
removeRdfStatementWithContextReturn	1
getPathwayAsResponse generateIDReturn	1
generateidketurn	
RunBlastnJobResponse	
indexWithSwAReturn	
GetPresentFamiliesHogenomResponse getSpeciesScientificNameReturn	
getspeciesscientificivameretum getresultReturn	
adjacentReturn	1 1
runDSSPResponse	1
listPathwaysForEntityIdsReturn	1 1
getSGDAccessionNumberReturn	Ī
sources	1
is archaealReturn	1
runProqDefaultResponse	1
getMappingsReturn getBlocksReturn	
getBlocksReturn	
getAllGeneReturn	1
getTermMetadataResponse	1
abstract coupled toReturn	<u>1</u>
runPfragDefaultResponse	<u>1</u>
fastaStructureIdQueryReturn getPBfromGOReturn	
uniprot acc	1
getFeaturesResponse	1 1
smilesQueryReturn	1
getSimpleModelsByReactomeldsReturn	1
getSimpleModelsByReactomeldsReturn getAllModelsReturn	Ī
UniProtId2DomainIdsResponse	1
getObsoletePdbldsReturn	1
getObsoletePdbldsReturn getModelsIdByUniprotIdReturn	1
mot listResponse	1
getSymbolInfoFromSymbolResponse	
cla_queryResponse getSecretoryOrNonsecretoryProteinsForAGivenGenomeReturn	1
getSecretoryOrNonsecretoryProteinsForAGivenGenomeReturn	
mappingInfoResponse	
queryResponse getlalgorithmReturn	
cla fetch propResponse	
getTheProteinsThatAreInTheSameOrthoMCLClusterAsAGivenProteinReturn	1 1
GetSpeciesListResponse	1 1
getInstanceByldentifierResponse	1 1
getInstanceByldentifierResponse sliceChromosomeByGeneResponse	1 1
blastn BasicReturn	$\overline{1}$
cla dump propResponse	1
cla dump propResponse getPfamAAnnotationForAGivenProteinReturn	1
queryForExperimentReturn getSpellingSuggestionsReturn getAvailableTaxaResponse	1
getSpellingSuggestionsReturn	1
getAvailableTaxaResponse	1
getESTsWithAGivenESTUnisequenceReturn GetPublicationExperimentsResponse	
GetPublicationExperimentsResponse	1
uniquesubsReturn getTermTypesReturn doScoobyBySeqResponse	1
getTermTypesKeturn	
uoscoopybyseqkesponse	<u> </u>
whatWholemountRegionExpressGeneReturn	<u>1</u>
getGeneList1Response getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameOrthoMCLClustersAsProteinsWithAGivenGOAnnotationReturn	
bindndnaResponse	
getEntriesFieldUrlsResponse	
getEntriesrieidonskesponse getDatabaseToFieldsMapReturn	1
doc2locResponse	1
doc2locResponse DecoderResponse	1
AsyncSimpleSearchResponse	1
getOrganismsByQueryTermResponse	1
gg	

name	count
getELMResponse	1
soap find clashesResponse RetrieveCompleteScoresResponse MOBYSHoundLocusIDFromConservedDomainDbIDReturn	1
RetrieveCompleteScoresResponse MORVSHoundLocusIDFromConsorvedDomainDhIDPoturn	 1
removeRdfStatementReturn	+ 1
getCurationTagHistoryResponse	+ 1
getCurationTagHistoryResponse performGOAnalysisReturn runhogenomResponse RunBJastpJobResponse	$\frac{1}{1}$
runhogenomResponse	1
RunBlastpJobResponse	1
transformReturn	1
GetInheritedFamiliesHogenomResponse	$\frac{1}{1}$
selectMagemlReturn	1 1
getAnnotationDateReturn gashsupReturn	
cluster_by_bbhsReturn	+ +
runDSSPByIDResponse	- 1
generatePathwayDiagramInSVGReturn	1
getGeneOntologyBiologicalProcessReturn is_bacterialReturn	1
is_bacterialReturn	1
GetDatabankInfoResponse	1
ligandsmiles	1
getSubmissionWithoutImageReturn getTermXrefsResponse	1 1
pinned_region_dataReturn	+ +
runPalignSeqSeqDefaultResponse	+ +
fastaQueryReturn	+ +
getNodeAccessionReturn	1
runGlobPlotterResponse	1
getReleaseDatesReturn	1
getSimpleModelsRelatedWithChEBIReturn	1
rachanca traeRaturn	1
PdbSuperfamily2UniProtIdsResponse getDomainFragmentsReturn getModelsIdByUniprotIdsReturn	$\frac{1}{1}$
getDomainFragmentsKeturn	1
cla_listResponse	
identifyDatabasesReturn	+ +
RunDeterministicSimulatorResponse	1
getSymbolInfoFromSymbolAndTaxIdResponse	1
hit queryResponse	1
getSignalPPredictionForAGivenProteinReturn	1
numberConversionResponse	1
GetDatabasesResponse	1
seq_fetch_fastaResponse getMCLClustersThatContainProteinsOfAGivenGenomeReturn	$\frac{1}{1}$
GetEnsIdsByRegionResponse	+ 1
getInstancesWithPdbEntryAssignedResponse	- 1
upLoadGenBankLocalFileResponse	$\frac{1}{1}$
blastp BasicReturn state of the	$\overline{1}$
seg dump fastaResponse	1
getProteinsofAGivenGenomeWithAGivenPfamAnnotationReturn	1
queryForArraysReturn	1
getESTUnisequenceWithAGivenIdentifierReturn	$\frac{1}{1}$
GetPublicationStrainsResponse	<u> </u>
parsimonyReturn getSplSetIdReturn	1 1
getSignalpReturn	 1
whatVoxelRegionExpressGeneReturn	+ 1
getRepresentativeImageSeriesResponse	$\frac{1}{1}$
getAnnotationForProteinsOfAGivenGenomeThatAreInMCLClustersWithSecretoryOrNonsecretoryProteinsReturn	1
bindnrnaResponse	1
getDomainsReferencedInEntryResponse	1
getChipTypesReturn	1
Domains 2 svgResponse	1
GetAsyncSearchStatusResponse	
	1 1

	count
getGeneldsByQueryTermResponse	1
getELMByldentifierResponse soap match alignment with modelResponse	<u> </u>
updatePathwayResponse	+ +
getCogsConservedDomainIdsByLocusLinkIdReturn	<u> </u>
getCogsConservedDomainIdsByLocusLinkIdReturn addRdfReturn	$\bar{1}$
getCurationTagsResponse getAffyNormalizationMethodsReturn	1
getAffyNormalizationMethodsReturn	1
analyzeGenesReturn DunPlactyleh Response	1
RunBlastxJobResponse	1
match_summary GetInnovatedFamiliesHogenomResponse	- †
IOUPLVDAIADVEXOURPLUID	$\frac{1}{1}$
getSequenceTypeReturn getReportResponse cluster_by_simReturn	1
getReportResponse	1
cluster_by_simReturn	1
runMkDomResponse	<u></u>
generatePathwayDiagramInSVGForldReturn getGeneOntologyMolecularFunctionReturn	+
is eukaryoticReturn	
is eukaryoticReturn GetMetaDataResponse	1
screeningsearchreturn	$\frac{1}{1}$
getAllSubmissionsReturn	1
getOntologyNamesResponse	1
ăll_genomesReturn	1
runPalignSeqSeqResponse homologyReductionQueryReturn TFmodellerReturn	$\frac{1}{1}$
nomologyReductionQueryReturn International Programme	<u></u>
getNodeInfoReturn	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
PdhSuperfamily2KeggOrthologsResponse	1
getEcNumsForStructuresReturn	$\frac{1}{1}$
PdbSuperfamily2KeggOrthologsResponse getEcNumsForStructuresReturn getSimpleModelsByChEBIldsReturn	$\overline{1}$
response alignmentReturn	1
GetHogenomFamCompResponse	1
blastQueryXmlReturn	<u> </u>
getSimpleModelsByUniprotIdsReturn hit listResponse	<u></u>
identifySimpleSearchesReturn	1
StartDeterministicSimulatorResponse	<u> </u>
getSymbolInfoFromReferenceResponse	$\overline{1}$
hat queryResponse	1
getTheNumberOfParalogsForAllMCLClustersContainingProteinsOfAGivenGenomeReturn	1
runMutalyzerResponse	<u> 1</u>
SearchByMassResponse	<u> </u>
pickingResponse mot fetch fastaResponse	
getAllMCLClustersWithMoreThanAGivenPercentageOfProteinsOfAGivenGenomeReturn	1
GetExpressionDatasetsResponse	1
getInstancesByPubMedIdResponse	$\overline{1}$
ŭpLoadGenBańkRemoteFileResponse	1
translating Blast BasicReturn	1
mot_dump_fastaResponse	$\frac{1}{1}$
getTheNumberOfGenomesInMCLClustersThatContainProteinsOfAGivenGenomeAndProteinsOfEssentialYeastGenesReturn	├
deleteReturn getProprietaryInformationPoturn	
getProprietaryInformationReturn phyloML	+ +
getESTsRepresentingAGivenOpenReadingFrameReturn [7]	<u> </u>
GetPublicationModelTypesResponse	$\frac{1}{1}$
maximumlikelihoodReturn	$\bar{1}$
findRemappedReturn	1
getMsblastInfoReturn	1
getSubmissionAssayWlzReturn	1
geneSearch1Response	<u></u>
getAnnotationForProteinsOfAGivenGenomeThatAreInOrthoMCLClustersWithSecretoryOrNonsecretoryProteinsReturn	
	1

	count
disemblResponse CotEMPL Nucleatide Sequence - Remil Response	$\frac{1}{1}$
GetEMBLNucleotideSequence BsmlResponse getDatabasesWithDetailsReturn	1
EncoderResponse	1
GetAsyncSearchResultResponse getGeneldsByProteomeldResponse getELMsByTextSearchResponse	1
getGeneldsByProteomeldResponse	1
getELMsByTextSearchResponse	1
applyReturn findPathwaysRyXrefResponse	<u>_</u>
getCddConservedDomainIdsBvLocusLinkIdReturn	1
findPathwaysByXrefResponse getCddConservedDomainIdsByLocusLinkIdReturn runFetchDomResponse removeRdfReturn	$\bar{1}$
removeRdfReturn	1
findInteractionsResponse	1
getTwoColorBackgroundCorrectionMethodsReturn getGOViewHTMLReturn	<u></u>
RunTblastnJobResponse	1
search summary	$\bar{1}$
GetLostFamiliesHogenomResponse	1
queryDataByArrayIdReturn	1
getSequenceSourceReturn external_callsReturn	<u></u>
runZpredDefaultResponse	1
load Dathway Farld Daturn	$\frac{1}{1}$
getGeneOntologyCellularComponentReturn	1
getGeneOntologyCellularComponentReturn is_prokaryoticReturn GetIndicesResponse	1
GetIndicesResponse	<u>_</u>
dockingscore getSubmissionAssayImageReturn	<u>_</u>
getOntologyLoadDateResponse	1
genomesReturn	1
runPalignSeqSeqSSDefaultResponse	1
keywordQueryReturn	1
outseq getNodeStageReturn	<u></u>
KeggOrtholog2PdbSuperfamiliesResponse	<u>_</u>
getEcNumsReturn	$\frac{1}{1}$
helloBioModelsReturn	1
response_sequencesReturn	1
GetHogenomFamDefResponse blastStructureIdQueryXmlReturn	<u>_</u>
getModelsIdByUniprotReturn getModelsIdByUniprotReturn	1
hat listResponse	1
identifySimpleSearchInputParametersReturn	1
getSymbolInfoFromIHOPResponse	1
hot queryResponse	$\frac{1}{1}$
getTheNumberOfParalogsForAllOrthoMCLClustersContainingProteinsOfAGivenGenomeReturn transcriptInfoResponse	<u>_</u>
SearchByMass2Response	1
mappingResponse	$\frac{1}{1}$
seg fetch nrResponse	1
getÄllMCLClustersWithLessThanAGivenPercentageOfProteinsOfAGivenGenomeReturn	1
GetPathwayListResponse	<u>_</u>
request_origResponse updateCollectionResults	<u></u> 1
blastn AdvancedReturn	1
seg dump nrResponse	1
get The Number Of Genomes In MCLC lusters That Contain Proteins Of A Given Genome And Proteins Of Only Essential Yeast Genes Return	1
runQueryDefaultDbReturn	1
getESTUnisequencesRepresentingAGivenOpenReadingFrameReturn	<u></u>
getCategoriesResponse distanceReturn	<u></u>
getSourceTypesReturn	1
getMsTagInfoReturn	1
getStoreSizeResponse	1
	i

name	count
geneSearchResponse	1
getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameMCLClustersAsProteinsWithAGivenPSortPredictionReturn	11
iupredlResponse	
getDetailledNumberOfResultsResponse GetEMBLNucleotideSequence AGAVEResponse	
getLinksReturn	1
FillseaResnanse	1
FillseqResponse GetCompoundThumbnailResponse getGeneldsByProteomeldAndQueryTermResponse getELMsBySequenceMatchResponse	1
getGeneldsBvProteomeldAndOuervTermResponse	1
getELMsBvSeguenceMatchResponse	1
ConvertCopasiMLToSBMLResponse getColoredPathwayResponse MOBYSHoundLocusIDFromOMIMReturn	1
getColoredPathwayResponse	1
MOBYSHoundLocusIDFromOMIMReturn	1
runFetchProtResponse	1
selectQuerySerializedReturn	1
loginResponse	1
getTwoColorWithinArrayNormalizationMethodsReturn	1
getGOViewPDFReturn	1
RunTblastxJobResponse	
filter file	
GetRegainedFamiliesHogenomResponse	
GetStpGraphListResponse	
getTranscriptIdReturn function ofReturn	
antDINTSPacerdRyAccessionDespanse	
getPRINTSRecordByAccessionResponse loadPathwayForObjectReturn	1
aetAffyldByOrfReturn	1
getAffyIdByOrfReturn getTimedNodesReturn	+ 1
translation ofReturn	1
getSnpsBySwissProtIdReturn	1
getCogldReturn	
getAllTermsFromOntologyResponse	1
replacesReturn	1
runProdivtmhmmDefaultResponse	1
pubmedAbstractQueryReturn	1
getRootAccessionReturn	1
KeggOrtholog2DomainIdsResponse	1
getCifChainReturn	1
getAllModelsIdReturn	1
ResultMS	1
getPubMedFromPMIDResponse	
getSNPIdsReturn	
getModelsIdByGOIdReturn	
hot_listResponse evaluateSimpleSearchReturn	
getSymbolInteractionsFromSymbolResponse	
mom_gueryResponse	1
mom_queryResponse getEnzymesForAGivenProteinReturn	1
SearchByFormulaResponse	1
getImageResponse	1
mot fetch nrResponse	1
getTheProteinsThatAreInTheSameMCLClusterAsAGivenProteinReturn	1
GetEnsIdsByPathwayNameResponse	1
PhysicoChemProfile	
deleteCollectionResults	
blastp AdvancedReturn	
mot dump nrResponse	
getTheNumberOfGenomesInMCLClustersThatContainProteinsOfAGivenGenomeAndProteinsOfOnlyNonessentialYeastGenesReturn	
retriveDocReturn	
getRxConceptPropertiesReturn	1
sequenceML	1
getESTUnisequencesForAGivenGenomeReturn	
getNonOboOntologiesResponse	
helpResponseSoapMsg	

name	count
getMsFitInfoReturn	1
generateConceptsFromPubMedQueryResponse getGeneList2Response	
getGeneListzKesponse	1
getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameOrthoMCLClustersAsProteinsWithAGivenPSortPredictionReturn iupredsResponse	
nupreuskesponse gotEntry/FioldUrlsPosponso	
getEntryFieldUrlsResponse getField2TableAndColumnMapReturn	1
FormatnameRecnonce	+ 1
FormatnameResponse GetCompoundInfoResponse getGeneIdsByGenomeAcResponse getAllELMInstancesResponse ConvertSBMLToCopasiMLResponse getPathwayInfoResponse MOBYSHoundGOIDFrome	<u>†</u>
getGeneldsBvGenomeAcResponse	1
getAllELMInstancesResponse	1
ČonvertSBMLToCopasiMLResponse	1
getPathwayInfoResponse	1
MOBYSHoundGOIDFromLocusIDReturn	1
runreichrastakesponse	
addRdfWithContextReturn	1
getRecentChangesResponse	1
get i woloiorBetweenArrayNormalizationMetnodsKeturn	1
getRecentChangesResponse getTwoColorBetweenArrayNormalizationMethodsReturn analyseGenesPDFOutputReturn GetTransferedFamiliesHogenomResponse	1
Get Hansiereuraniines nogenomikes ponse Get Str. Granh Resnonse	
GetStpGraphResponse getTargetDescriptionReturn genomes ofReturn getPRINTSRecordByIDResponse listPathwayParticipantsReturn getInterproByOrfReturn getAbstractNodesReturn	1
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genomes oncean	1
listPathwayParticipantsReturn	+ 1
getInterproBvOrfReturn	1
getAbstractNodesReturn	1
uel translationnetum	1
CountResponse	1
getRootTermsResponse	1
taxonomy ofReturn runProdivtmhmmParamsResponse	1
runProdivtmhmmParamsResponse	
getSpaceGroupForStructureReturn	1
srs-result	
getOntologyReturn	1
DomainId2KeggOrthologsResponse getCifResidueReturn	
getClinesiddeReturn getAllCuratedModelsIdReturn	
eInfoResult	1
UniProt ID	+ 1
getPubMedFromIHOPResponse	1
getUniProtSNPsReturn	1
getModelsIdByGOReturn	Ī
seq_infoResponse	1
identifyAdvancedSearchCollectionsReturn	1
getSymbolInteractionsFromSymbolAndTaxIdResponse	1
seq_fetch_nameResponse	1
getProteinsofAGivenGenomeWithAGivenEnzymeAnnotationReturn	1
addResult	1
SearchByFormula2Response	
getKinaseListResponse	1
seq_dump_nameResponse	1
getTheNumberOfProteinsForEachGenomeInAGivenMCLClusterReturn GotEnsIdsByExpressionBosponso	<u>1</u>
ĞetEnsIdsByExpressionResponse Clu	<u>1</u>
ciu updateResourceResults	1
translating Blast AdvancedReturn	1
hit fetch simpleResponse	1
getTheNumberOfGenomesInOrthoMCLClustersThatContainProteinsOfAGivenGenomeAndProteinsOfEssentialYeastGenesReturn	+ 1
retriveDocDefaultDhReturn	+ †
getRelatedByRelationshipReturn	<u>† †</u>
getRelatedByRelationshipReturn contactResponse	<u>† 1</u>
geneSearchReturn	1
getESTsForAGivenGenomeReturn	1

name	count
getQualityCriteriaResponse	
submit jobResponseSoapMsg getTicketByIdResponse	
gerneketbyldkesponse generateConceptsFromTextResponse	
runAtlasResponse	
getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameMCLClustersAsProteinsWithAGivenWolfPSortPredictionReturn	
pdisorderResponse	
listFieldsResponse]
getLocationResponse]
annotateProbesReturn	
PhylotableResponse Mol2CSIDResponse getGeneByldResponse getELMInstanceResponse	
moizCSIDRESpoise gotGonoByldPosnonso	
getGenebylukesponse netFLMInstanceResponse	
ValidateModelResponse	
listPathwaysResponse	
listPathwaysResponse MOBYSHoundLocusIDFromGOIDAndECodeReturn	7
runFetchAlianResponse	
addRdfFileWithContextReturn	
getPathwayResponse	
getGOViewImageReturn	
getDescriptionReturn	
NearestNodeProdomResponse UploadStpGraphResponse getArchivalUniGeneClusterReturn	
optoaustpuraprikesponse actArchivalUniConoClustorPoturn	
gerarchivatoriideneciusternetum genus speciesReturn	
genus speciesReturn getSeedAlignmentByAccessionResponse listPathwayParticipantsForIdReturn	
listPathwayParticipantsForldReturn	
patchGetGeneSymbolReturn	
getSliceReturn	
feature locationReturn]
CooccurrenceResponse]
getAnnotationsReturn	
getTermsByNameResponse	
search_and_grepReturn	
runTopcompResponse getPubmedIdForAllStructuresReturn	
getPcaAnalysisReturn	
getSupportedLibsReturn	
getSupportedElbSNeturn getOntologyTermReturn	
getOntologyTermReturn PdbSuperfamily2KeggPathwaysResponse	7
getChainsReturn	
getAllNonCuratedModelsIdReturn]
eInfoResultMS]
FASTA	
getRelatedSymbolsResponse	
getNCBIdbSNPDescriptorReturn getModelsIdByTaxonomyIdReturn	
not_infoResponse	
dentifyAdvancedSearchFiltersReturn	
getSymbolInteractionsFromReferenceResponse	+ =
not fetch nameResponse	
getProteinsofAGivenGenomeWithTheSameEnzymeAnnotationAsAGivenProteinReturn	
odeSolverResult	
GetRecordMolResponse	
getBindingDomainListResponse	
not dump nameResponse	
getASummaryOfTheMCLClusterAnalysisForAGivenGenomeReturn	
GetExpressionByEnsIdsResponse Cl2	
appendResourceResults	
getTagldentInfoReturn	
nat fetch simpleResponse	
getTheNumberOfGenomesInOrthoMCLClustersThatContainProteinsOfAGivenGenomeAndProteinsOfOnlyEssentialYeastGenesReturn	+ =

name	coun
GetExperimentResponse	
addDocumentReturn	
getRelatedByTypeReturn getPipelinesStatusResponse	 ;
searchReturn	
retCellularI ocalisationPredictionPSortWolfPSortForAGivenProteinReturn	
getResourcesByCategoryResponse istDomainsResponse getTicketBySeqResponse generateTerminologyResponse etchAtlasResultResponse	
istDomainsResnonse	
retTicketBvSeaResnonse	
gerneketbysegnesponse generateTerminologyResponse	
etchAtlasResultResponse	
JelannolationForProteinsUtaGivenGenome InatareinTheSameUmoWCLCUSterSASProteinSWithAGivenWoltPSortPredictionReturn	
poodlelResponse getEntriesResponse getGeneldResponse	-
getEntriesResponse	
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dsToProbesReturn	
PsiBlast NTResponse	
CSID2ExtRefsResponse	
getAllFunctionalSitesResponse	
learReturn	
findPathwaysByLiteratureResponse MOBYSHoundGiFromLocusIDReturn	
MOBYSHoundGiFromLocusIDReturn	
runFetchKWResponse removeRdfWithContextReturn MUSCLEAlignment	
removeRdfWithContextReturn	
MUSCLEAlignment	
performJustAffyNormalizationReturn	;
getBiologicalProcessGOTermByOrfReturn GetAvailableDatabasesResponse	;
jetAvailabieDatabaseskesponse	
resourcePathReturn	
GetParentProdomResponse CYGDAnnotations	 ;
getUniGeneldReturn	
id2dnaReturn	
getSeedAlignmentByIDResponse	
naxSizeInListObjects	
patchGetTranscriptIdReturn	
getWholemount2DInfoReturn	
nastructAlignmentML	
ooundaries ofReturn	-
SpellCheckResponse	-
getEntitiesReturn	
getSegldReturn	
getTermsByExactNameResponse	
simple_searchReturn	
getStructureGenomicsPdbldsReturn	
somReturn	
ArrayOfOperonResult	
getTimedNodeAccessionReturn KeggPathway2PdbSuperfamiliesResponse getCifChainsReturn	
KeggPathway2PdbSuperfamiliesResponse	;
getUitChainsReturn	;
getModelByIdReturn	;
eSearchResult	;
queueentry rotPolatodSymbolsWithTaxIdPospopso	
jetRelatedSymbolsWithTaxIdResponse getModelsIdByTaxonomyReturn	
cla_infoResponse	 :
dentifyAdvancedSearchFilterOperatorsReturn	
getSynsetsReturn	
getSynsetsReturn getSymbolInteractionsFromIHOPResponse	
cla_fetch_nameResponse	
getProteinsofAGivenGenomeInAGivenPathwayReturn	
nathPlotResult	
GetCompressedRecordsSdfResponse	
getSubstratesByNameSearchResponse	
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ament Response in Committee Committ	name	count
MCLC Users That Contain Proteins Ol AGroup Ol Genomes Return	nfo	1
DEGEniedResponse DeResponse	cla_dump_nameResponse	1
Jake Resurre Alexable Jake Alexable Jake Resurre Alexable Jake Re	getMCLClustersThatContainProteinsOfAGroupOfGenomesReturn	1
Interest	AffyDEGFinderResponse	
tick stamplescaponas (citch simplescaponas (citch simplescaponas (citch simplescaponas (citch simplescaponas (citch simplescaponas (citch simplescaponas)) (citch simplescaponas) (citc	MSÍ	$\frac{1}{1}$
fects smipleResponse InhelimberOfechomesinOrthoMCLClustersThatContainProteinsOfAGivenGenomeAndProteinsOfOnlyNonessentialYeastGenesReturn InhelimberOfechomesinOrthoMCLClustersThatContainProteinsOfAGivenGenomeAndProteinsOfOnlyNonessentialYeastGenesReturn IDDccumentDefaultDbReturn IDDccumentDbReturn IDDCcumentDefaultDbReturn IDDCcumentDbReturn IDDCcumentDbBReturn IDDCcume		<u>l</u>
The Number Of Genomes In Orthon MCL Clusters That Contain Proteins Of Adversing Of Only Nonessential Yeast Genes Return Experiments Response All Related in Cartery of the Contain Proteins Of Adversing Of Only Nonessential Yeast Genes Return Ty Prindides poors Submission Annotation ExtReturn Submission Annotation ExtReturn Ty Proteins of Adversing Annotation Cartery of Contain Proteins of Adversing Annotation Return WSDLS Response Low State of Contain Contain Contain Proteins Of Adversing Annotation For Treatment of Contain Contai		<u>l</u>
ExperimentsResponse	10t_fetch_simplekesponse rotThoNumberOfConomosInOrthoMCl ClustersThotContainDrotainsOfACivenConomoAndDrotainsOfOnlyNenescentialVeastConosDaturn	<u></u>
DocumentDefaultDeReturn	Set Fynoriments Posponso	<u></u>
AllRelatedInfoReturn Vymindlesponse bitoni retReturn InchlasonReturn Inchlason	addDocumentDefaultDhReturn	1
styPmidResponse 1 1 1 1 1 1 1 1 1	retΔIIRelatedInfoReturn	+ 1
SubmissionAnnotation PextReturn ChapsonReturn ChapsonRetur		$\frac{1}{1}$
rchjasonReturn Trotte insoft Asiven Genome With AGI ven Cellular Localisation As Predicted By PSort Return 1	netSubmissionAnnotationTextReturn	1
ProteinsofAcivenGenomeWithAcivenCellularLocalisationAsPredictedByPSortReturn DomainsReterenceInDomainResponse 1	searchlason Return	1
DomainsketereneedInthomainkesponse	petProteinsofAGivenGenomeWithAGivenCellularLocalisationAsPredictedBvPSortReturn	$\frac{1}{1}$
DomainsketereneedInthomainkesponse	netWSDLsResponse	1
Intelligeny Return 1 1 1 1 1 1 1 1 1	getDomainsReferencedInDomainResponse	1
AnnotationForProteins/0fAgrenGenomeThatAreInTheSameMCLClustersAsProteinsWithAgivenEnzymeAnnotationReturn TeleferencedEntilesSeponse	pronetOuervReturn	${1}$
AnnotationForProteins/0fAgrenGenomeThatAreInTheSameMCLClustersAsProteinsWithAgivenEnzymeAnnotationReturn TeleferencedEntilesSeponse	plastprodRésponse	1
Indexesponse	getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameMCLClustersAsProteinsWithAGivenEnzymeAnnotationReturn	1
ReferencedEntriesResponse	nahindrResponse	1
TunctionalsiteResponse	getReferencedEntriesResponse	
TunctionalsiteResponse	getGeneResponse	1
TunctionalsiteResponse	keywordsToProbesReturn	1
TunctionalsiteResponse	PsiBlast_NT_XMLResponse	1
1 10 10 10 10 10 10 10	getFunctionalSiteResponse	1
1	getRepositoriesReturn	1
FetchKWscriptResponse	removeCurationTagResponse	1
CallsweaveReturn	MOBYSHoundLocusIDFromGiListReturn	<u>l</u>
CallsweaveReturn	TunFetchKWscriptResponse	<u>l</u>
GenebescriptionByOrfReturn	emoveRatfileWithcontextReturn	<u>l</u>
InputDescriptionReturn 1 1 1 1 1 1 1 1 1	/OIDCAIISWEAVERETURN	<u>_</u>
ChildProdomResponse Spy QueryReturn Subsystem ClassificationsReturn Subsystem ClassificationsReturn Spy QueryReturn Spy QueryReturn Spy QueryReturn Spy Spy	getGeneDescriptionByOnReturn	<u></u>
SyQueryReturn	JetinputDescriptionReturn CotChildProdomPocnopso	<u></u>
GenomeVersionReturn 1 1 1 1 1 1 1 1 1	istRyOugh/Poturn	<u></u>
2dnaReturn		- 1 1
PIRSeedAlignmentByAccessionResponse	ide?dnaRaturn	1
KeggPathwayASGifReturn 1 ChGeTargetDescriptionReturn 1 Wholemount2DReturn 1 PSFIlesReturn 1 gestSearchTermsResponse 1 GlimmersReturn 1 PrefixedTermsByNameResponse 1 subsystem classificationsReturn 1 ApropertyFetchResultResponse 1 SwissProtldsReturn 1 ayofRegulatorResult 1 AllStageNameReturn 1 igPathway2DomainIdsResponse 1 ChainLengthReturn 1 tarchResultMS 1 SymbolsFromReferenceResponse 1 SymbolsFromReferenceResponse 1 SubModelSBMLReturn 1 infoResponse 1 uluateAdvancedSearchReturn 1 UniprotiDReturn 1 LatestSymbolinformationFromSymbolResponse 1	retPIRSeedAlianmentRvAccessionResnonse	1
1	get inseedalighmentbyAccessionnesponse netKengPathwayΔcGifReturn	1
PSFilesReturn 1 gestSearchTermsResponse 1 GlimmersReturn 1 PrefixedTermsByNameResponse 1 subsystem classificationsReturn 1 ApropertyFetchResultResponse 1 SwissProtidsReturn 1 ayOfRegulatorResult 1 AllStageNameReturn 1 gpathway2DomainIdsResponse 1 ChainLengthReturn 1 ModelSBMLByIdReturn 1 surchResultMS 1 SymbolsFromReferenceResponse 1 SubModelSBMLReturn 1 infoResponse 1 luateAdvancedSearchReturn 1 UniprotIDReturn 1 LatestSymbolInformationFromSymbolResponse 1	patchGetTargetDescriptionReturn	1
PSFilesReturn 1 gestSearchTermsResponse 1 GlimmersReturn 1 PrefixedTermsByNameResponse 1 subsystem classificationsReturn 1 ApropertyFetchResultResponse 1 SwissProtidsReturn 1 ayOfRegulatorResult 1 AllStageNameReturn 1 gpathway2DomainIdsResponse 1 ChainLengthReturn 1 ModelSBMLByIdReturn 1 surchResultMS 1 SymbolsFromReferenceResponse 1 SubModelSBMLReturn 1 infoResponse 1 luateAdvancedSearchReturn 1 UniprotIDReturn 1 LatestSymbolInformationFromSymbolResponse 1	netWholemount2DReturn	1
1 1 1 1 1 1 1 1 1 1	netPSFilesReturn	1 1
ggestSearchTermsResponse1GlimmersReturn1PrefixedTermsByNameResponse1subsystem classificationsReturn1ApropertyFetchResultResponse1SwissProtIdsReturn1ayOfRegulatorResult1AllStageNameReturn1igPathway2DomainIdsResponse1ChainLengthReturn1ModelSBMLByIdReturn1barchResultMS1SymbolsFromReferenceResponse1SubModelSBMLReturn1infoResponse1IutateAdvancedSearchReturn1UniprotIDReturn1LatestSymbolInformationFromSymbolResponse1		1
1 1 1 1 1 1 1 1 1 1	SuggestSearchTermsResponse	1
PrefixedTermsByNameResponse subsystem classificationsReturn ApropertyFetchResultResponse SwissProtldsReturn SwissProtldsReturn 12 3QOfRegulatorResult 13 3QOfRegulatorResult 14 3QOfRegulatorResult 15 3QOFRegulatorResult 16 3QOFRegulatorResult 17 3QOFRegulatorResult 18 3QOFRegulatorResult 19 3QOFReg	getGlimmersReturn	1
subsystem classificationsReturn ApropertyFetchResultResponse SwissProtIdsReturn ayOfRegulatorResult AllStageNameReturn gipPathway2DomainIdsResponse ChainLengthReturn IndelSBMLByIdReturn IndelSBMLByIdReturn IndelSBMLByIdReturn InfoResponse		1
SwissProtIdsReturn 1 ayOfRegulatorResult 1 AllStageNameReturn 1 gPathway2DomainIdsResponse 1 ChainLengthReturn 1 ModelSBMLByIdReturn 1 sarchResultMS 1 SymbolsFromReferenceResponse 1 SubModelSBMLReturn 1 infoResponse 1 luteteAdvancedSearchReturn 1 UniprotIDReturn 1 LatestSymbolinformationFromSymbolResponse 1	all subsystem classificationsReturn	1
ayOfRegulatorResult AllStageNameReturn ggPathway2DomainIdsResponse ChainLengthReturn ModelSBMLByIdReturn earchResultMS SymbolsFromReferenceResponse SubModelSBMLReturn 11 SubModelSBMLReturn 12 SubModelSBMLReturn 13 SubModelSBMLReturn 14 SubModelSBMLReturn 15 SubModelSBMLReturn 16 SubModelSBMLReturn 17 SubModelSBMLReturn 18 SubMod	DNApropertyFetchResultResponse	1
AllStageNameReturn ggPathway2DomainIdsResponse ChainLengthReturn ModelSBMLByldReturn earchResultMS SymbolsFromReferenceResponse SubModelSBMLReturn infoResponse 1 luateAdvancedSearchReturn UniprotIDReturn 1 LatestSymbolInformationFromSymbolResponse	getSwissProtIdsReturn	1
AllStageNameReturn ggPathway2DomainIdsResponse ChainLengthReturn ModelSBMLByldReturn earchResultMS SymbolsFromReferenceResponse SubModelSBMLReturn infoResponse 1 luateAdvancedSearchReturn UniprotIDReturn 1 LatestSymbolInformationFromSymbolResponse	ArrayOfRegulatorResult	1
ModelSBMLByIdReturn EarchResultMS SymbolsFromReferenceResponse SubModelSBMLReturn InfoResponse IlluateAdvancedSearchReturn	getAllStageNameReturn	1
ModelSBMLByIdReturn EarchResultMS SymbolsFromReferenceResponse SubModelSBMLReturn InfoResponse IlluateAdvancedSearchReturn	KeggPathway2DomainIdsResponse	1
ModelSBMLByIdReturn EarchResultMS SymbolsFromReferenceResponse SubModelSBMLReturn InfoResponse IlluateAdvancedSearchReturn	getChainLengthReturn	1
SymbolsFromReferenceResponse SubModelSBMLReturn infoResponse IluateAdvancedSearchReturn UniprotIDReturn LatestSymbolInformationFromSymbolResponse	getModelSBMLByldReturn	1
SubModelSBMLReturn1infoResponse1iluateAdvancedSearchReturn1UniprotIDReturn1LatestSymbolInformationFromSymbolResponse1	eSearchKesultMS	
infoResponse iluateAdvancedSearchReturn UniprotIDReturn 1 LatestSymbolInformationFromSymbolResponse	getSymbolsFromKeterenceResponse	
lluateAdvancedSearchReturn UniprotIDReturn 1 LatestSymbolInformationFromSymbolResponse	getSubModelSBMLKeturn	
UniprotIDReturn LatestSymbolInformationFromSymbolResponse 1		
LatestSymbolInformationFromSymbolResponse 1		
LatestSymbolinformationFromSymbolkesponse		$\frac{1}{2}$
TOTON MODELLACINONO	getLatestSymbolinforMationFromSymbolKesponse	<u> </u>
_ietcn_desckesponse1	seq_fetch_descResponse	

name	count
getProteinsofAGivenGenomeThatAreInTheSamePathwayAsAGivenProteinReturn	1
prismResult CotResponde Off Response	<u></u>
GetRecordsSdfResponse getSubstratesByGeneNameSearchResponse	<u></u>
seq_dump_descResponse	1
getTheGeneForAGivenGeneSymbolReturn	1
getDatabaseVersionResponse	+ 1
CoiledCoils	1
searchResults	1
tblastnReturn	$\frac{1}{1}$
guery listResponse	1
getMCLClustersContainingProteinsOfAGivenGenomeAndProteinsOfEssentialYeastGenesReturn	1
GetExperimentsBySearchResponse	1
deleteDefaultDbReturn	1
getMultiIngredBrandReturn	1
getSubmissionAnnotationWholemountWlzReturn	1
searchMFqueryReturn	1
getProteinsofÁGivenGenomeWithAGivenCellularLocalisationAsPredictedByWolfPSortReturn	
ÄfoldSOAPResponse	<u> </u>
getReferencedEntriesSetResponse	<u>_</u>
blastprodout isServiceAliveReturn	<u></u>
getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameOrthoMCLClustersAsProteinsWithAGivenEnzymeAnnotationReturn	<u>L</u>
say_helloResponse	<u> </u>
get Organism Response	- 1
getOrganismResponse PsiBlast_ProteinResponse	1
toAlignedDotBracketFastaReturn	+ 1
getFunctionalSitesByTextSearchResponse	1
selectQueryReturn	1
listOrganismsResponse	$\frac{1}{1}$
MOBYSHoundGiFromLocusIDListReturn	$\frac{1}{1}$
getUPIForSequenceResponse	1
indexFromRemoteReturn	1
addStepAffyNormalizationReturn	1
getOrfByGenenameReturn	1
getMethodExampleReturn	1
GetDFStreeProdomResponse	1
listObjectsReturn	1
getAlignmentsReturn	<u>l</u>
contigs_ofReturn	<u></u>
getPIRSeedAlignmentByIDResponse	<u></u>
getKeggldsByKeggPathwayReturn getOrfsByGoldReturn getStagesWithReconstructionReturn	<u></u>
getOTSDyGOTUREtuTT	<u></u>
getDotFileReturn	1
pegs_with_md5_stringReturn	<u></u> †
BlastResponse	1
getIdStatusReturn	+ 1
getTermParentsResponse	1
all familiesReturn	$\frac{1}{1}$
trnascanFetchResultResponse	1
getGenomeDetailsReturn	1
ÄrrayOfProfileParameterResult	1
getStageNameReturn	1
DomainId2KeggPathwaysResponse	1
DomainId2KeggPathwaysResponse getCifChainLengthReturn getSimpleModelByIdReturn eSummaryResult	1
getSimpleModelByldReturn	1
eSummaryResult	1
guessSymbolidFromSymboliextResponse	1
hat infoResponse	1
getESTLibraryWithAGivenIdentifierReturn getESTLibraryWithAGivenIdentifierReturn	
ČheckSyntaxŘesponse	$\frac{1}{2}$
getLatestSymbolInformationFromSymbolAndTaxIdResponse	
mot_fetch_descResponse	
	I

name	count
getPathwaysForAGivenProteinReturn	1
asyncResult	$\overline{1}$
GetExtendedCompoundInfoResponse	1
getInstancesByProteinSequenceAccessionSearchWithAllThroughputDataResponse	1
mot dump descResponse	1
getTheProteinsForAGivenGeneReturn getTheProteinsForAGivenGeneReturn	1
getSourceDatabaseVersionsResponse	1
manovaReturn	1
getSourcesResults	1
tblastxReturn	1
query_infoResponse	1
getMCLClustersContainingProteinsOfAGivenGenomeAndProteinsOfOnlyEssentialYeastGenesReturn	1
GetExperimentPhenoAssaysResponse	1
createDbReturn	1
getDrugsReturn	1
getSubmissionAnnotationVoxelWlzReturn	1
getProteinsofAGivenGenomeWithTheSamePredictedPSortCellularLocalisationAsAGivenProteinReturn	1
ContrafoldSOAPResponse	1
listFieldsInformationResponse	1
blastpgpout	1
getAnnotationForProteinsOfAGivenGenomeThatAreInTheSameMCLClustersAsProteinsInAGivenPathwayReturn	
topconsResponse	
getProteinResponse	$\frac{1}{1}$
PsiBlast_Protein_XMLResponse	
fromAlignedDotBracketFastaReturn	
getELMInstanceBySequenceMatchResponse	1
addRdfFileReturn	$\frac{1}{1}$
getCurationTagsByNameResponse	<u> </u>
listFilesReturn	1
getUPIForAccessionResponse	$\frac{1}{1}$
indexFromSRBReturn	
performJustTwoColorPreprocessingReturn performJustTwoColorPreprocessingReturn	1
getSGDÁccessionByOrfReturn getInputExampleReturn	$\frac{1}{1}$
getinputExampleKeturn	
GetPresentFamiliesProdomResponse	
listTopLevelPathwaysReturn	
getGeneTitleReturn	1 11