The EMERALD ontology efforts

James Malone (malone@ebi.ac.uk)

European Bioinformatics Institute. Wellcome Trust Genome Campus, Hinxton, Cambridge, CB10 1SD.

Fundamental the successful analysis and reproducibility of microarray to experiments is the quality of the documentation and descriptions that are used to report microarray experiments. lt is because of the value of MIAME (Minimum standardised reporting that initiatives such as Information About а Microarray Experiment) and MAGE-TAB (MicroArray Gene Expression Tabular) emerged rapidly adopted by the microarray and were Recently, towards а more formalised representation community. а movement the knowledge in form of ontologies become increasingly of has community. widespread biomedical likes of OBO within the The the (Open Biomedical Ontologies) Foundry have helped to enable those wishing to knowledge representation techniques model their use formal to data using ontology standards such as OBO and OWL (Web Ontology Language). advantages Ontologies offer the of shared understanding, reducina ambiguity, richer machine representations of data and degree of а readability, allowing computation to be performed over such models such towards as consistency checking. We discuss а move improving elements of the reporting of microarray experiments with the development of an and ontology for Normalisation Data Transformation. We also highlight the contribution from the OBI (Ontology **Biomedical** Investigations) for efforts and this reporting experiments Consortium how may impact the of within the microarray community...