|  |  |  |  |
| --- | --- | --- | --- |
| Classifier group | Classifier | Relevance assignment | Example |
|  |  |  |  |
| MAIN CLASSIFIERS | Data type | **High**: abundance, density, EBV genetic analysis.  **Moderate**: distribution, presence-absence.  **Low**: presence-only, relative abundance, species richness, non-EBV genetic analysis. | L |
| Spatial range | **High**: >15.000 km2.  **Moderate**: 5.000 to 15.000 km2.  **Low**: <5.000 km2. | M |
| Temporal range | **High**: >10 years.  **Moderate**: 2 to 10 years.  **Low**: <1 to 2 years. | M |
| **Assigned relevance by Main Classifiers**  (criteria: majority vote) | | | **M** |
|  |  |  |  |
| MODULATORS | Multispecies | >10 species. | FALSE |
| Threatened species | Assigned a threatened level by IUCN. | TRUE |
| New species to science | A first description of a species. | FALSE |
| New regional species | A first record of a species in the region. | FALSE |
| Priority area\* | Data from a geographic area of special interest. | FALSE |
| **Assigned relevance by Main Classifiers + Modulators**  **(**criteria: if any TRUE, relevance by Main Classifiers increases one category) | | | **H** |
| \*For our study case, the ATLAS database, we prioritize areas from Northern Quebec. | | | |