

Readme Data Endopaths

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Link to git: https://gitlab.engine.capgemini.com/ti/france/data-and-ai/dimedia/endopath_new

This is the list of all the data that are important or might be useful, or that are used in the pipeline. There are numerous other data versions whose purpose is not clear and that can be ignored.

| Name | Explanation | Path |
|---------------------------------------|--|---------------------------------|
| Raw | | |
| dossier-gyneco-23-03-2022.csv | Raw, unprocessed data with medical texts. Opening this file requires a password which can be found in 'Raw' > 'Nouveau dossier' > 'PASSWORD_1.txt' | Raw |
| Recueil (1).csv | Data with demographical informations and symptoms, '1' specifying presence, '0' absence of symptom. Used in the Machine Learning scripts to predict endometriosis. | Raw |
| Recueil_final.csv | Like 'Receuil (1)', but with tab that explains the meaning of the different columns. Not used in any script or notebook. | Raw |
| Processed | | |
| all_words.txt | All words that appear in the text. Created in 'create_correct_dict_nlp.ipynb' | DATA_PROCESSED /Correction_mots |
| comparaison_gynéco_receuil.csv | This file was created manually. For each feature, the present/absent values are noted for the original recueil data on the one hand and the infos extracted manually from the gyneco files on the other. This is to compare the two and resolve conflicting informations. Lines that are marked in red present cases where the two files seem to give conflicting information (to be reviewed by a medical expert). | DATA_PROCESSED |
| data_gynéco_manual_extraction_raw.csv | This was generated by going through the gynéco files and | DATA_PROCESSED |

| | | |
|-----------------------------------|---|------------------------------------|
| | noting all the informations manually. It is not the final file version used in the code because it contains some comments etc. | |
| data_gynéco_manual_extraction.csv | Notes all the informations (present/absent) for the features as they have been found in the gynéco files by manual inspection. In difference to 'data_gynéco_manual_extraction_raw', the column names have the same format as in the recueil data here, and everything is cleanly formatted. Not finalized yet as of October '24. Copy missing columns in question from 'comparison_gynéco_recueil.csv'. | DATA_PROCESSED |
| data_synth_priorité_gynéco.csv | Data synthesized between the gynéco and recueil data. In cases of conflicting information between the two (the cases marked in red in 'comparison_gynéco_recueil.csv'), prioritizes the gynéco data. | DATA_PROCESSED |
| data_synth_priorité_recueil.csv | Data synthesized between the gynéco and recueil data. In cases of conflicting information between the two (the cases marked in red in 'comparison_gynéco_recueil.csv'), prioritizes the recueil data. | DATA_PROCESSED |
| dictionnaire_correction.json | Dictionary of misspelled words and their corrections for later use. Created in 'create_correct_dict_nlp.ipynb' | DATA_PROCESSED /Correction_mots |
| donnees_entree_nlp_sans_endo.csv | Data with medical texts, preprocessed and corrected, without the target columns referring to endometriosis. Used for NLP scripts. | DATA_PROCESSED |

The NLP/ML scripts in many cases have variables in the beginning of the script by which you can choose which type of data you want to enter into the model (e.g. either the original recueil data or the informations extracted manually from gynéco files). The documentation on Endopaths normally mentions in the top right of the slide which data have been used.