```
/* Takes study data ({@link mrmc.core.InputFile#keyedData}, {@link mrmc.core.InputFile#truthVals})
 *  and creates data for {@link mrmc.core.CovMRMC}
* --t-matrices: reader scores
* ---- {@link #t0_modAA}, {@link #t0_modAB}, {@link #t0_modBB}: signal-absent scores  [Nnormal ][Nreader][2 modalities]
* ---- {@link #t1_modAA}, {@link #t1_modAB}, {@link #t1_modBB}: signal-present scores [Ndisease][Nreader][2 modalities]
* --d-matrices: study design
* ---- {@link #d0_modAA}, {@link #d0_modAB}, {@link #d0_modBB}: signal absent indicator [Nnormal ][Nreader][2 modalities]
* ---- {@link #d1_modAA}, {@link #d1_modAB}, {@link #d1_modBB}: signal present scores   [Ndisease][Nreader][2 modalities]
* --fully crossed status
* ---- {@link #fullyCrossedA}, {@link #fullyCrossedB}, {@link #fullyCrossedAB}
*
* CALLED BY: {@link mrmc.gui.InputFileCard.varAnalysisListener} */
```