Appendix C

Formal Description of Scenario 1

 ${f section}\ Framework\ {f parents}\ standard_toolkit$

```
[CHAR, DATE, DECISION]
LIGAND == \operatorname{seq} CHAR
RECEPTOR == \operatorname{seq} CHAR
CONFIG == \operatorname{seq} CHAR
RESULT == \operatorname{seq} CHAR
LIGANDS == \mathbb{P} LIGAND
RECEPTORS == \mathbb{P} RECEPTOR
RESULTS == \mathbb{P} RESULT
DEEP\_ALIGN\_RESULT == \operatorname{seq} CHAR
YES\_NO ::= yes \mid no
USER\_INPUT == \operatorname{seq} CHAR
PREVIOUS\_RESULT == (LIGAND \times RECEPTOR \times CONFIG \times DATE) \mapsto RESULT
PREVIOUS\_RESULTS == \{PREVIOUS\_RESULT\}
```

```
\begin{array}{c} \textit{dockingWithConfig}: (\textit{LIGAND} \times \textit{RECEPTOR} \times \textit{CONFIG}) \rightarrowtail \textit{RESULT} \\ \hline \\ \forall \textit{l}: \textit{LIGAND}; \; \textit{r}: \textit{RECEPTOR} \mid \textit{l} \neq \varnothing \land \textit{r} \neq \varnothing \bullet \exists \textit{c}: \textit{CONFIG}; \; \textit{res}: \textit{RESULT} \mid \\ \textit{c} \neq \varnothing \bullet \textit{dockingWithConfig}(\textit{l},\textit{r},\textit{c}) = \textit{res} \\ \end{array}
```

 $Docking_AutoDockVina$

ligand?: LIGAND

receptor?: RECEPTOR

config?: CONFIG result!: RESULT

 $config? \neq \emptyset \land result! = docking With Config (ligand?, receptor?, config?)$

 $Molecular Docking Environment_Raccoon 2$ _

ligands?:LIGANDS

receptors?: RECEPTORS

config?: CONFIG results!: RESULTS

date!: DATE

 $\exists \ ligand?: ligands?; \ receptor?: receptors?; \ result!: results! \bullet Docking_AutoDockVina$

 $. \ View Molecular Docking Results_Raccoon 2$

 $\Xi Molecular Docking Environment_Raccoon2$

 $results! \neq \emptyset$

 $. \ Molecular Docking Results Repository _MongoDB \ __$

 $repository: (\mathit{LIGAND} \times \mathit{RECEPTOR} \times \mathit{CONFIG} \times \mathit{DATE}) \leftrightarrow \mathit{RESULT}$

 $decisionRepository: \{PREVIOUS_RESULT\} \leftrightarrow DECISION$

 $repository \neq \emptyset$

```
 LinsertUpdateMolecularDockingResultsRepository1 $$ \Delta MolecularDockingResultsRepository\_MongoDB $$ l?: LIGAND $$ r?: RECEPTOR $$ c?: CONFIG $$ res?: RESULT $$ d?: DATE $$ repository' = repository <math>\oplus \{(l?, r?, c?, d?) \mapsto res?\} $$
```

```
Select Molecular Docking Results
\Xi Molecular Docking Results Repository\_MongoDB
whereL?:LIGAND
where R?: RECEPTOR
where C?: CONFIG
where D?: DATE
where Res?: RESULT
selectResults!: (LIGAND \times RECEPTOR \times CONFIG \times DATE) \leftrightarrow RESULT
liq: LIGAND
rec: RECEPTOR
con: CONFIG
dat: DATE
selectResults! = \{(where L?, where R?, where C?, where D?)\} \triangleleft repository \lor
selectResults! = \{(where L?, where R?, where C?, dat)\} \triangleleft repository \lor
selectResults! = \{(where L?, where R?, con, where D?)\} \triangleleft repository \lor
selectResults! = \{(where L?, where R?, con, dat)\} \triangleleft repository \lor
selectResults! = \{(where L?, rec, where C?, where D?)\} \lhd repository \lor \\
selectResults! = \{(where L?, rec, where C?, dat)\} \triangleleft repository \lor
selectResults! = \{(where L?, rec, con, where D?)\} \triangleleft repository \lor
selectResults! = \{(where L?, rec, con, dat)\} \triangleleft repository \lor
selectResults! = \{(liq, where R?, where C?, where D?)\} \triangleleft repository \lor
selectResults! = \{(lig, where R?, where C?, dat)\} \triangleleft repository \vee
selectResults! = \{(liq, where R?, con, where D?)\} \triangleleft repository \lor
selectResults! = \{(lig, where R?, con, dat)\} \triangleleft repository \lor
selectResults! = \{(lig, rec, where C?, where D?)\} \triangleleft repository \lor
selectResults! = \{(lig, rec, where C?, dat)\} \lhd repository \lor
selectResults! = \{(lig, rec, con, where D?)\} \triangleleft repository \lor
selectResults! = \{(lig, rec, con, dat)\} \triangleleft repository \lor
 selectResults! = repository \triangleright \{whereRes?\}
```

```
\forall \ dar : DEEP\_ALIGN\_RESULT; \ ui : USER\_INPUT \bullet \exists \ threshold : \mathbb{Z}; \ DeepScore : \mathbb{Z} \bullet DeepScore \geq threshold \wedge goodDeepAlignResult(dar, ui) = yes \vee DeepScore < threshold \wedge goodDeepAlignResult(dar, ui) = no
\_AssessDeepAlign\_
```

 $goodDeepAlignResult: (DEEP_ALIGN_RESULT \times USER_INPUT) \leftrightarrow YES_NO$

 $DeepAlignResult?: DEEP_ALIGN_RESULT$ $userInput?: USER_INPUT$ r: RECEPTOR $assessed_receptors!: RECEPTORS$ $r \in assessed_receptors! \land goodDeepAlignResult(DeepAlignResult?, userInput?) = yes \lor$

 $r \not\in assessed_receptors! \land goodDeepAlignResult(DeepAlignResult?, userInput?) = no$

```
SelectMolecularDockingResults \\ previous\_results?: PREVIOUS\_RESULTS \\ userInput?: USER\_INPUT \\ assessedPreviousResults!: PREVIOUS\_RESULTS \\ \\ \exists previous\_result: previous\_results?; result: RESULT \bullet \\ \{result\} = \operatorname{ran}(previous\_result) \land \\ (previous\_result \in assessedPreviousResults! \land goodDocking(result, userInput?) = yes \lor \\ previous\_result \not\in assessedPreviousResults! \land goodDocking(result, userInput?) = no) \\ \end{cases}
```

```
 \begin{aligned} make A \, Decision A \, dditional \, Tool : & (RECEPTORS \times PREVIOUS\_RESULTS) \\ \leftrightarrow & DECISION \\ \\ \hline \exists \, previous\_results : PREVIOUS\_RESULTS; \, receptors : RECEPTORS; \\ lig : LIGAND; \, d : DECISION; \, con : CONFIG; \, dat : DATE \bullet \\ \forall \, previous\_result : previous\_results; \, receptor : receptors \bullet \\ \{(lig, receptor, con, dat)\} = \operatorname{dom}(previous\_result) \\ \land \\ make A \, Decision A \, dditional \, Tool(receptors, previous\_results) = d \end{aligned}
```

```
\_DecisionMaker\_Custom = \\ assessed\_previous\_results?: PREVIOUS\_RESULTS \\ assessed\_receptors?: RECEPTORS \\ decision!: DECISION \\ \hline \\ decision! = \\ makeADecisionAdditionalTool(assessed\_receptors?, assessed\_previous\_results?)
```