

The new fragment picker

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RosettaCON 2010

Summary

- 1 Why fragmentpicker?
- 2 Fragmentpicker overview
- 3 Fragmentpicker quick examples
 - NNmake - style protocol
 - CS-Rosetta type protocol
- 4 Fragmentpicker details
 - All about quota
 - Fragments' scoring system
- 5 Fragmentpicker: under the hood
- 6 The end
 - There is no end...
 - Acknowledgements

When three is not enough...

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- `core/fragment/picking/*` by Andrew Ban
(→ `core/fragment/picking_old/`)

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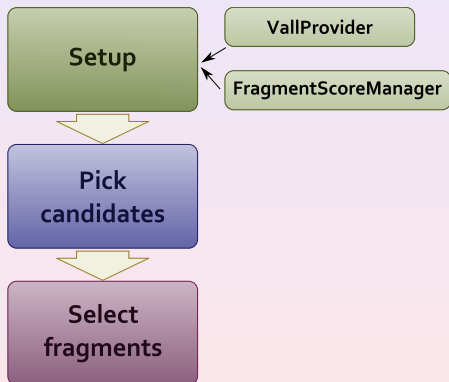
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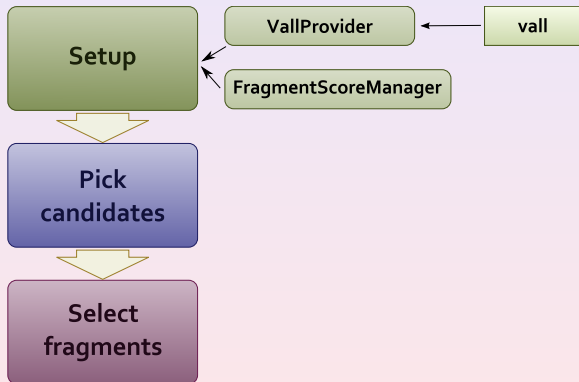
Overview & input files



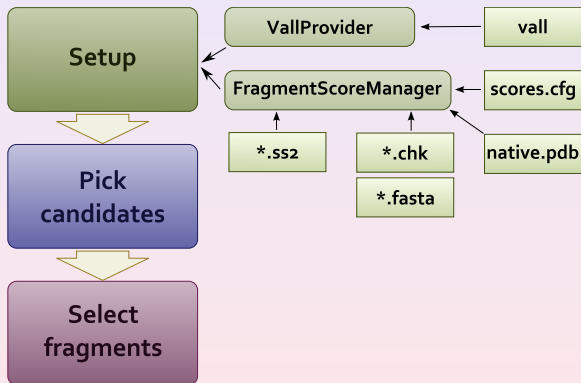
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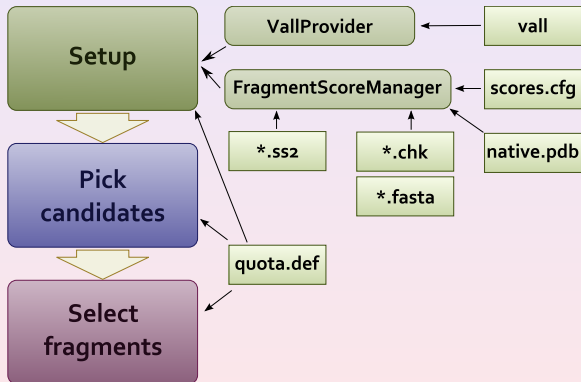
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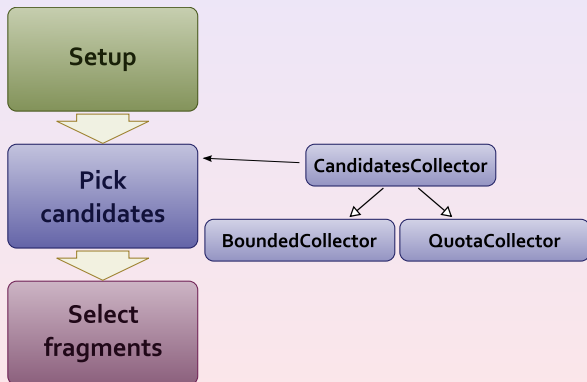
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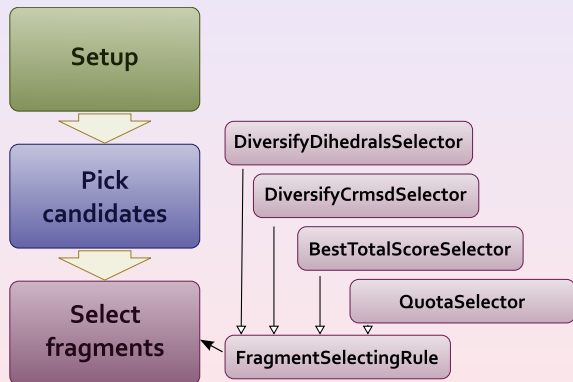
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NNmake - style protocol

See Psi-Jufo-Sam-L1-Quota demo

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- scoring:(file: psi_sam_jufo_L1-Q.cfg)

```
# score name          priority wght cutoff extras
SecondarySimilarity   350 1.0 - psipred
SecondarySimilarity   300 1.0 - sam
SecondarySimilarity   250 1.0 - jufo
RamaScore              150 50.0 -
ProfileScoreL1         200 1.0 -
PhiPsiSquareWell       100 0.0 -
FragmentCrmsd          30 0.0 -
```

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```

- quota setup (file: Quota.def):

```
#pool_id pool_name fraction
1  psipred  0.6
2  jufo     0.2
3  sam      0.2
```


CS-Rosetta protocol

See CS-Talos-L1-Rama demo

- collector: BoundedCollector

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#	score	name	priority	wght	cutoff	extras
	CSScore		375	3.0	-	
	RamaScore		400	50.0	-	talos
	SecondarySimilarity		350	3.0	-	talos
	ProfileScoreL1		200	1.0	-	
	PhiPsiSquareWell		100	0.0	-	
	FragmentCrmsd		30	0.0	-	
	GunnCostScore		20	0.0	-	

Picking results

- fragment files

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- fragment files
- score files

How quota works

Vall fragment

12asA	F	E	167	163	...
12asA	V	E	168	162	...
12asA	H	E	169	161	...
12asA	S	H	170	160	...
12asA	Q	H	171	159	...
12asA	E	H	172	158	...
12asA	L	H	173	157	...
12asA	L	H	174	156	...
12asA	S	H	175	155	...

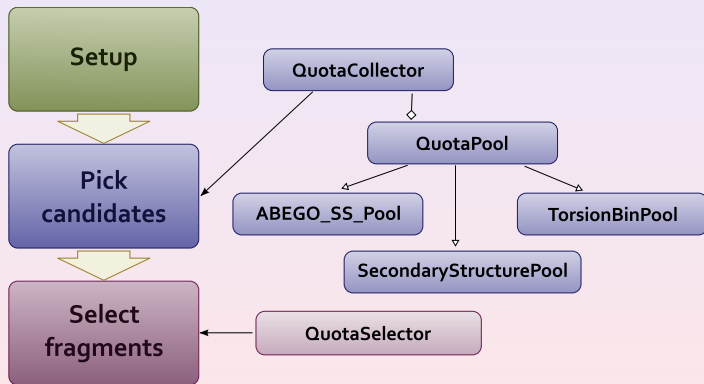
PsiPred prediction

16	T	E	0.030	0.016	0.924
17	T	E	0.065	0.207	0.636
18	T	E	0.101	0.258	0.569
19	E	H	0.114	0.586	0.348
20	A	H	0.083	0.335	0.582
21	V	H	0.125	0.521	0.354
22	D	H	0.545	0.177	0.215
23	A	H	0.079	0.916	0.011
24	A	H	0.052	0.933	0.003

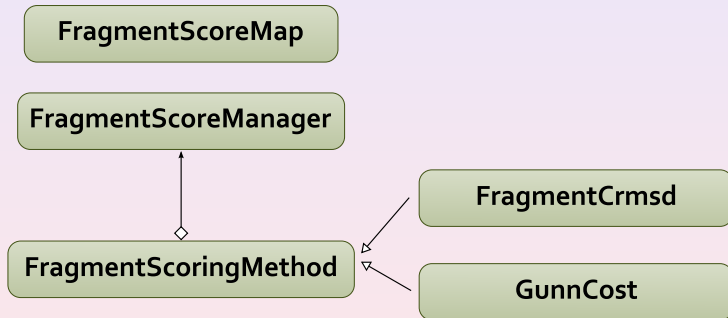
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3 sam 0.2
```

```
psipred-C : 0.125 × 200 × 0.6 = 15
psipred-H : 0.521 × 200 × 0.6 = 63
psipred-E : 0.354 × 200 × 0.6 = 42
```

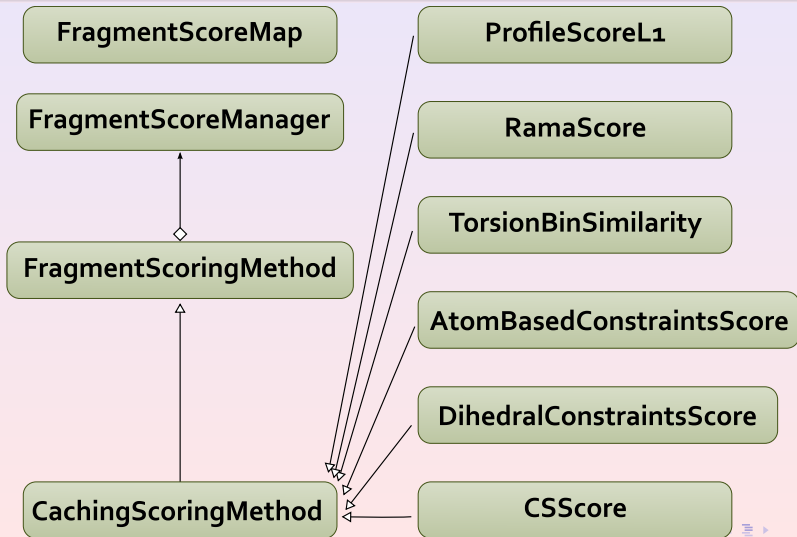
Quota-related parts



Scoring-related classes



Scoring-related classes (2)



fragmentpicker_integration_demo.cc

see

[mini/src/apps/pilot/dgront/fragmentpicker/fragmentpicker_integration_demo.c](#)

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- pick any fragment size you want: `-frags::frag_sizes 3 6 9 17`
- restrict sequence range: `-frags::picking::query_pos 110 111`
- drink Porter, no more Jufo

Future plans

- more scores

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- more experimental data

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- SQL database instead of vall

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