Report on Identified Refactoring Opportunities for (SC/ST) Refactoring

The tool identifies the Control-Fields and conditional constructs (Switch/If) that use these Control-Fields to simulate the (SC / ST) refactoring. It prioritizes the Control-Fields (refactoring opportunities) based on the following criteria:

- Number of conditional constructs that switch on Control-Field
 Group i represents i different conditional constructs where the Control-Field is used
- 2. Average size of the conditional body
- 3. Number of control values
 - 2-3 control values
 - 3-6 control values
 - 6-n control values
- 4. Presence of conditional constructs with respect to the class of declaration (COD) of Control-Field
 - In COD (A)
 - Outside (B)
 - Mixed (C)
- 5. Qualified for SC or ST
- 6. Static Field
- 7. Have subclasses already

Input Benchmark Statistics:

No. Of Classes: 102

No. Of Primitive Felds: 312

No. Of Control-Fields for Subclass Pattern: 5 No. Of Control-Fields for State pattern: 12

Total No. Of Control-Fields: 17

Uses	Replace Type Code with Subclass (SC)	
5	<pre><graphics.readviz, forwarddirection=""></graphics.readviz,></pre>	(A)
3	<misc.alignmentblock, strand=""></misc.alignmentblock,>	(A)
	<pre><graphics.readviz.readmapcomparat forward="" or,=""></graphics.readviz.readmapcomparat></pre>	(A)
2	<pre><rnaseq.finesplicecounter.splicelocati exon1="" on,=""></rnaseq.finesplicecounter.splicelocati></pre>	(A)
	<misc.filter.junction, top=""></misc.filter.junction,>	(A)

Uses	Replace Type Code with State (ST)	
9	<misc.alignmentrecord, forwardStrand></misc.alignmentrecord, 	(C)
8	<pre><misc.filter.filterinterfaceadaptor,< td=""><td>(B)</td></misc.filter.filterinterfaceadaptor,<></pre>	(B)
4	<pre><rnaseq.alignmentfilter, quiet=""></rnaseq.alignmentfilter,></pre>	(A)
3	<misc.alignmentfilter2, quiet=""></misc.alignmentfilter2,>	(A)
	<pre><rnaseq.exoncounter, introniccgff=""></rnaseq.exoncounter,></pre>	(C)
	<pre><rnaseq.alignmentfilter, forcesamout=""></rnaseq.alignmentfilter,></pre>	(A)
	<misc.alignmentfilter2, isProcessNonPassed></misc.alignmentfilter2, 	(A)
	<pre><graphics.readviz, inauto=""></graphics.readviz,></pre>	(A)
2	<pre><special.intronretentioncgff2,< td=""><td>(A)</td></special.intronretentioncgff2,<></pre>	(A)
	<pre><rnaseq.transcriptomerecover, intronfiltermode=""></rnaseq.transcriptomerecover,></pre>	(A)
	<pre><misc.filter.filterinterfacesinglereada< td=""><td>(C)</td></misc.filter.filterinterfacesinglereada<></pre>	(C)
	<pre><special.promotercgff, fromtss=""></special.promotercgff,></pre>	(A)

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