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Figure. 4. RESCUE-ESEs and PESEs in sequences found by SR protein responsive functional SELEX (Liu et al. 2000; Liu et al. 1998). The ~34-mers shown include a central 20 nt derived from a random insert, plus constant flanks of 7 nt each; the 20-mers represent the sequences underlying ESEfinder (Cartegni et al. 2003), kindly provided by Adrian Krainer. RESCUE-ESEs are in bold blue, PESEs are in bold red, overlaps between the two are in violet italics and the top ESEfinder sequences (largest increment over the default threshold) are underlined (thin = ASF/SF2, thick = SC35, dotted = SRp40, and dashed = SRp55). Turn off spell check for better viewing in Microsoft Word.