		Window Sizes (base pairs)																				
		10	20	50	80	100	200	500	800	1000	2000	5000	8000	10000	20000	50000	80000	100000	200000	500000	800000	1E+06
Human Genome	chr1	0.637	0.756	0.883	0.928	0.945	0.978	0.993	0.996	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr2	0.626	0.749	0.879	0.924	0.942	0.976	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr3	0.613	0.738	0.873	0.920	0.939	0.975	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr4	0.610	0.736	0.872	0.920	0.938	0.975	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr5	0.616	0.741	0.874	0.921	0.939	0.975	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000
	chr6	0.618	0.742	0.875	0.921	0.939	0.975	0.992	0.995	0.996	0.997	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000
	chr7	0.637	0.757	0.884	0.928	0.944	0.978	0.993	0.996	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr8	0.623	0.746	0.877	0.923	0.941	0.976	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr9	0.639	0.759	0.885	0.929	0.945	0.978	0.993	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr10	0.634	0.755	0.881	0.926	0.943	0.977	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000
	chr11	0.635	0.756	0.884	0.928	0.945	0.978	0.993	0.996	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr12	0.628	0.749	0.878	0.924	0.942	0.977	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000
	chr13	0.619	0.744	0.876	0.922	0.940	0.976	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr14	0.635	0.755	0.883	0.928	0.944	0.978	0.993	0.996	0.997	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr15	0.636	0.756	0.882	0.926	0.943	0.977	0.993	0.995	0.996	0.998	0.999	0.999	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000
	chr16	0.646	0.761	0.884	0.928	0.945	0.978	0.993	0.995	0.996	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000
	chr17	0.657	0.770	0.889	0.931	0.947	0.979	0.993	0.996	0.996	0.998	0.999	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000
	chr18	0.622	0.746	0.876	0.922	0.940	0.975	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000
	chr19	0.660	0.769	0.888	0.930	0.947	0.979	0.992	0.995	0.995	0.997	0.998	0.998	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000
	chr20	0.638	0.756	0.882	0.926	0.943	0.977	0.992	0.995	0.996	0.998	0.999	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr21	0.653	0.772	0.893	0.934	0.949	0.980	0.994	0.996	0.997	0.998	0.999	0.999	0.999	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000
	chr22	0.658	0.770	0.888	0.929	0.946	0.978	0.992	0.995	0.996	0.997	0.998	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999	0.999
	chrX	0.598	0.724	0.864	0.915	0.935	0.973	0.991	0.994	0.995	0.997	0.998	0.998	0.999	0.999	0.999	0.999	0.999	0.999	1.000	1.000	1.000
	chrY	0.602	0.728	0.867	0.917	0.936	0.973	0.990	0.993	0.994	0.996	0.997	0.998	0.998	0.998	0.998	0.998	0.999	0.998	0.999	0.998	0.998

		Window Sizes (base pairs)																				
		10	20	50	80	100	200	500	800	1000	2000	5000	8000	10000	20000	50000	80000	100000	200000	500000	800000	1E+06
Randomized Human Genome	chr1	0.366	0.488	0.671	0.759	0.796	0.881	0.936	0.949	0.954	0.966	0.972	0.974	0.975	0.976	0.977	0.976	0.977	0.977	0.978	0.977	0.976
	chr2	0.365	0.487	0.670	0.758	0.795	0.882	0.936	0.950	0.955	0.966	0.973	0.975	0.975	0.977	0.978	0.978	0.978	0.976	0.977	0.979	0.977
	chr3	0.365	0.487	0.669	0.758	0.795	0.882	0.937	0.951	0.956	0.966	0.973	0.975	0.975	0.976	0.975	0.976	0.976	0.976	0.977	0.976	0.979
	chr4	0.363	0.485	0.668	0.757	0.794	0.881	0.937	0.951	0.955	0.967	0.973	0.975	0.975	0.976	0.977	0.977	0.977	0.978	0.977	0.981	0.982
	chr5	0.365	0.487	0.670	0.758	0.795	0.881	0.937	0.951	0.955	0.967	0.973	0.975	0.975	0.977	0.979	0.979	0.979	0.981	0.979	0.981	0.977
	chr6	0.365	0.487	0.670	0.758	0.795	0.882	0.937	0.950	0.955	0.967	0.973	0.975	0.976	0.977	0.977	0.978	0.977	0.977	0.976	0.978	0.978
	chr7	0.365	0.487	0.669	0.758	0.795	0.881	0.937	0.950	0.955	0.966	0.973	0.975	0.976	0.978	0.978	0.978	0.979	0.979	0.979	0.981	0.981
	chr8	0.365	0.487	0.670	0.758	0.796	0.882	0.937	0.951	0.955	0.967	0.973	0.975	0.976	0.977	0.978	0.978	0.979	0.980	0.982	0.982	0.983
	chr9	0.366	0.488	0.670	0.759	0.796	0.881	0.935	0.948	0.951	0.966	0.972	0.974	0.975	0.976	0.977	0.974	0.976	0.976	0.974	0.973	0.973
	chr10	0.366	0.488	0.671	0.759	0.795	0.881	0.936	0.950	0.954	0.966	0.972	0.974	0.975	0.976	0.977	0.978	0.979	0.979	0.974	0.977	0.975
	chr11	0.366	0.488	0.671	0.759	0.795	0.881	0.936	0.950	0.954	0.966	0.973	0.974	0.975	0.976	0.976	0.977	0.977	0.977	0.977	0.980	0.980
	chr12	0.366	0.488	0.670	0.758	0.796	0.882	0.937	0.950	0.955	0.966	0.972	0.974	0.975	0.977	0.978	0.978	0.978	0.977	0.976	0.973	0.976
	chr13	0.363	0.485	0.668	0.757	0.795	0.881	0.937	0.950	0.954	0.967	0.974	0.976	0.977	0.978	0.979	0.978	0.978	0.979	0.974	0.981	0.981
	chr14	0.365	0.487	0.670	0.759	0.796	0.882	0.937	0.951	0.956	0.966	0.973	0.975	0.976	0.977	0.977	0.977	0.977	0.976	0.980	0.978	0.978
	chr15	0.366	0.488	0.670	0.759	0.795	0.881	0.936	0.949	0.954	0.966	0.973	0.974	0.976	0.977	0.978	0.979	0.978	0.978	0.976	0.978	0.975
	chr16	0.367	0.489	0.671	0.759	0.796	0.881	0.935	0.949	0.954	0.964	0.971	0.973	0.974	0.975	0.975	0.977	0.976	0.977	0.976	0.978	0.982
	chr17	0.366	0.488	0.670	0.758	0.794	0.880	0.934	0.948	0.952	0.964	0.970	0.972	0.972	0.973	0.975	0.975	0.974	0.971	0.974	0.971	0.973
	chr18	0.365	0.487	0.670	0.758	0.795	0.881	0.937	0.951	0.956	0.967	0.974	0.976	0.976	0.978	0.979	0.978	0.978	0.980	0.979	0.979	0.973
	chr19	0.367	0.489	0.671	0.758	0.794	0.879	0.933	0.947	0.949	0.962	0.968	0.971	0.971	0.972	0.972	0.970	0.966	0.965	0.968	0.971	0.970
	chr20	0.367	0.489	0.671	0.759	0.796	0.881	0.934	0.950	0.955	0.965	0.971	0.973	0.974	0.975	0.975	0.976	0.972	0.972	0.973	0.967	0.975
	chr21	0.366	0.488	0.671	0.759	0.796	0.881	0.936	0.949	0.953	0.965	0.972	0.973	0.974	0.975	0.976	0.976	0.976	0.970	0.968	0.959	0.960
	chr22	0.367	0.489	0.671	0.758	0.794	0.879	0.932	0.948	0.947	0.962	0.969	0.971	0.972	0.975	0.976	0.978	0.977	0.977	0.985	0.981	0.985
	chrX	0.365	0.486	0.669	0.758	0.795	0.881	0.936	0.950	0.953	0.966	0.973	0.975	0.975	0.976	0.977	0.978	0.978	0.980	0.975	0.975	0.979
	chrY	0.365	0.487	0.670	0.758	0.796	0.881	0.934	0.947	0.946	0.967	0.974	0.975	0.976	0.977	0.978	0.975	0.977	0.969	0.976	0.977	0.970