

<b>Code</b>	<b>MONTE_CARLO_SIMULATION.PY</b>
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<b>Summary</b>	Uses Monte Carlo simulation to test the central limit theorem, by generating and averaging random draws from several constituent distributions (triangular).
<b>Methods/ process</b>	<p><a href="#">Monte Carlo simulation</a>:</p> <ul style="list-style-type: none"> <li>- Method for evaluating complex probability spaces, where the output depends on several constituent probability distributions.</li> <li>- Can evaluate probabilistic quantities for which closed-form analytic solutions do not exist.</li> <li>- Constituent distributions can be of any type/family, discrete or continuous.</li> </ul> <p><a href="#">Central limit theorem</a>:</p> <ul style="list-style-type: none"> <li>- States that the average value of a sequence of random variables, each independently sampled, is normally distributed.</li> <li>- Asymptotically correct, as the number of distributions (and observations) increases.</li> <li>- Holds regardless of the types/parameters of the constituent distributions.</li> </ul> <p>Steps:</p> <ol style="list-style-type: none"> <li>1. Define the constituent distributions.</li> <li>2. Generate one random sample from each of those distributions (sampling them independently, with replacement).</li> <li>3. Compute the average value of those random samples (unweighted).</li> <li>4. Repeat this process for many sets of random draws (enough to sufficiently cover the sample space and generate a reasonably stable output distribution).</li> <li>5. Assess the distribution of the average value across all trials.</li> </ol>
<b>Training data</b>	<p>Constituent distributions:</p> <ul style="list-style-type: none"> <li>- 10 <a href="#">triangular</a> distributions (each with three parameters: minimum, mode, and maximum).</li> <li>- Fixed-width, non-overlapping, and abutting (no space between).</li> <li>- Mode value alternating between minimum and maximum values.</li> <li>- Each distribution is highly non-normal (skewed, with linear density segments).</li> <li>- Resembles a sawtooth wave, when all of the distributions are viewed together.</li> <li>- 1 million observations randomly generated from each distribution.</li> </ul>
<b>Output</b>	<p>Plots:</p> <ul style="list-style-type: none"> <li>- Scatter matrix (univariate histograms, 2D relationships)</li> <li>- Constituent distributions (sawtooth-like shape)</li> <li>- Average value distribution (histogram, cumulative probability)</li> </ul>
<b>Result</b>	Although the constituent distributions are highly non-normal, the distribution of their average value is a smooth bell-shaped curve, as the central limit theorem predicts.