

Brunner-Munzel検定について

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文献

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1 準備

1.1 パッケージの読み込みなど

```
In [1]: 1 using Base.Threads
2 using BenchmarkTools
3 using Distributions
4 using PrettyPrinting
5 using QuadGK
6 using Random
7 using RCall
8 using Roots
9 using StatsBase
10 using StatsFuns
11 using StatsPlots
12 default(fmt=:png, size=(400, 250),
13         titlefontsize=10, guidefontsize=8, tickfontsize=6)
14
15 x ≈ y = x < y || x ≈ y
16 x ≈̄ y = x > y || x ≈ y
17 safemul(x, y) = x == 0 ? x : isinf(x) ? typeof(x)(Inf) : x*y
18 safediv(x, y) = x == 0 ? x : isinf(y) ? zero(y) : x/y
```

Out[1]: safediv (generic function with 1 method)

1.2 組み合わせの生成子

In [2]:

```
1 """
2     nextcombination!(n, t, c = typeof(t)[min(t-1, i) for i in 1:t])
3
4 '[1,2,...,n]' からの重複無しの 't' 個の組み合わせ 'c' をすべて生成したい。
5
6 'nextcombination!(n, t, c)' は配列で表現された組み合わせ 'c' をその次の組み合わせに書き換えて,
7
8 初期条件を 'c = typeof(t)[min(t-1, i) for i in 1:t]' にすると, 'binomial(n, t)' 回の 'nextcomb
9 """
10 function nextcombination!(n, t, c = typeof(t)[min(t-1, i) for i in 1:t])
11     t == 0 && return c
12     @inbounds for i in t:-1:1
13         c[i] += 1
14         c[i] > (n - (t - i)) && continue
15         for j in i+1:t
16             c[j] = c[j-1] + 1
17         end
18         break
19     end
20     c
21 end
22 """
23 """
24 mycombinations!(n::Integer, t, c)
25
26 事前に割り当てられた組み合わせを格納する配列 'c' を使って, '[1,2,...,n]' からの重複無しの 't' 個
27 """
28 function mycombinations!(n::Integer, t, c)
29     for i in 1:t c[i] = min(t - 1, i) end
30     (nextcombination!(n, t, c) for _ in 1:binomial(n, t))
31 end
32 """
33 """
34 mycombinations!(a, t, c)
35
36 事前に割り当てられた組み合わせを格納する配列 'c' を使って, 配列 'a' からのインデックスに重複が
37 """
38 function mycombinations!(a, t, c)
39     t < 0 && (t = length(a) + 1)
40     (view(a, indices) for indices in mycombinations!(length(a), t, c))
41 end
42 """
43 """
44 mycombinations(x, t)
45
46 'x' が整数ならば '[1,2,...,x]' からの, 'x' が配列ならば 'x' からのインデックスに重複がない 't' 个
47 """
48 mycombinations(x, t) = mycombinations!(x, t, Vector{typeof(t)}(undef, t))
```

Out[2]: mycombinations

1.3 Welchのt検定

```
In [3]: 1 function tvalue_welch(m, x̄, sx², n, ȳ, sy²; Δμ=0)
2     (x̄ - ȳ - Δμ) / √(sx²/m + sy²/n)
3 end
4
5 function tvalue_welch(x, y; Δμ=0)
6     m, x̄, sx² = length(x), mean(x), var(x)
7     n, ȳ, sy² = length(y), mean(y), var(y)
8     tvalue_welch(m, x̄, sx², n, ȳ, sy²; Δμ)
9 end
10
11 function degree_of_freedom_welch(m, sx², n, sy²)
12     (sx²/m + sy²/n)² / ((sx²/m)²/(m-1) + (sy²/n)²/(n-1))
13 end
14
15 function degree_of_freedom_welch(x, y)
16     m, sx² = length(x), var(x)
17     n, sy² = length(y), var(y)
18     degree_of_freedom_welch(m, sx², n, sy²)
19 end
20
21 function pvalue_welch(m, x̄, sx², n, ȳ, sy²; Δμ=0)
22     t = tvalue_welch(m, x̄, sx², n, ȳ, sy²; Δμ)
23     v = degree_of_freedom_welch(m, sx², n, sy²)
24     2ccdf(TDist(v), abs(t))
25 end
26
27 function pvalue_welch(x, y; Δμ=0)
28     m, x̄, sx² = length(x), mean(x), var(x)
29     n, ȳ, sy² = length(y), mean(y), var(y)
30     pvalue_welch(m, x̄, sx², n, ȳ, sy²; Δμ)
31 end
32
33 function confint_welch(m, x̄, sx², n, ȳ, sy²; α=0.05)
34     v = degree_of_freedom_welch(m, sx², n, sy²)
35     c = quantile(TDist(v), 1-α/2)
36     SEhat = √(sx²/m + sy²/n)
37     [x̄-ȳ-c*SEhat, x̄-ȳ+c*SEhat]
38 end
39
40 function confint_welch(x, y; α=0.05)
41     m, x̄, sx² = length(x), mean(x), var(x)
42     n, ȳ, sy² = length(y), mean(y), var(y)
43     confint_welch(m, x̄, sx², n, ȳ, sy²; α)
44 end
```

Out[3]: confint_welch (generic function with 2 methods)

1.4 単峰型の函数が正の値になる場所を見つける函数

```
In [4]: 1 function findpositive(f, a, b; maxsplit = 30)
2     @assert f(a) < 0
3     @assert f(b) < 0
4     c = (a + b)/2
5     f(c) > 0 && return c
6     w = b - a
7     for k in 2:maxsplit
8         for d in range(w/2^(k+1), w/2-w/2^(k+1), step=w/2^k)
9             x = c + d
10            f(x) > 0 && return x
11            x = c - d
12            f(x) > 0 && return x
13        end
14    end
15    error("k > maxsplit = $maxsplit")
16 end
17
18 f(x) = abs(x) < 1e-4 ? 1.0 : -1.0
19
20 @time findpositive(f, -100abs(randn()), 20abs(randn()))
```

0.000092 seconds

Out[4]: -2.0987373957837008e-5

1.5 2つの分布が「互角」になるシフトの仕方を求める函数

In [5]:

```
1     """
2         prob_x_le_y(distx::UnivariateDistribution, disty::UnivariateDistribution;
3             a = 0.0)
4
5     この 函数は、連続分布 `distx`、`disty` と実数 `a` について、
6     `distx` と `disty` に従って生成される乱数をそれぞれ X, Y と書くとき、
7      $X \leq Y + a$  が成立する確率を返す。
8     """
9     function prob_x_le_y(distx::UnivariateDistribution, disty::UnivariateDistribution,
10         a = 0.0)
11         H(y) = cdf(distx, y) * pdf(disty, y-a)
12         quadgk(H, extrema(disty + a)...)[1]
13     end
14
15     """
16     tieshift(distx::UnivariateDistribution, disty::UnivariateDistribution;
17             p = 0.5)
18
19     この 函数は、連続分布 `distx`、`disty` と実数 `p` について、
20     `distx` と `disty` に従って生成される乱数をそれぞれ X, Y と書くとき、
21      $X \leq Y + a$  が成立する確率が `p` に等しくなるような実数 a を返す。
22     """
23     function tieshift(distx::UnivariateDistribution, disty::UnivariateDistribution;
24             p=0.5)
25         find_zero(a → prob_x_le_y(distx, disty, a) - p, 0.0)
26     end
27
28     @show tieshift(Normal(0, 1), Normal(2, 2))
29     @show tieshift(Normal(0, 1), Laplace(2, 2))
30     @show tieshift(Normal(0, 1), Uniform(0, 1));

```

```
tieshift(Normal(0, 1), Normal(2, 2)) = -1.999999999999232
tieshift(Normal(0, 1), Laplace(2, 2)) = -1.9999999999994498
tieshift(Normal(0, 1), Uniform(0, 1)) = -0.4999999999999983
```

2 Brunner-Munzel検定

2.1 Brunner-Munzel検定の実装

In [6]:

```
1 """
2     h_brunner_munzel(x, y)
3
4 この函数は, x < y のとき 1.0 を, x = y のとき 0.5 を返す.
5 """
6 h_brunner_munzel(x, y) = (x < y) + (x == y)/2
7
8 @doc raw"""
9     phat_brunner_munzel(X, Y)
10
11 まず以下のようにおく:
12
13 ````math
14 \begin{aligned}
15 &
16 H(x, y) = \begin{cases} 1 & (x < y) \\ 1/2 & (x = y) \end{cases}, \end{aligned}
17 \\
18 m = \mathrm{length}(X), \quad
19 n = \mathrm{length}(Y), \quad
20 x_i = X[i], \quad
21 y_j = Y[j]
22 \end{aligned}
23 ````

24 この函数は次の  $\hat{p}$  を返す:
25
26 ````math
27 \hat{p} = \frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n H(x_i, y_j).
28 ````

29 """
30 phat_brunner_munzel(X, Y) = mean(h_brunner_munzel(x, y) for x in X, y in Y)
31
32 @doc raw"""
33     statistics_brunner_munzel(X, Y,
34         Hx = similar(X, Float64),
35         Hy = similar(Y, Float64);
36         p = 1/2
37     )
38
39 この函数はデータ 'X', 'Y' について, Brunner-Munzel検定関係の統計量達を計算する. 詳細は以下の通り.
40
41 函数  $H(x, y)$  と  $\hat{p}$ ,  $H^x_i$ ,  $H^y_j$ ,  $\bar{H}^x$ ,  $\bar{H}^y$  を次のように定める:
42
43 ````math
44 \begin{aligned}
45 &
46 m = \mathrm{length}(X), \quad
47 n = \mathrm{length}(Y), \quad
48 x_i = X[i], \quad
49 y_j = Y[j],
50 \\
51 &
52 \hat{p} = \frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n H(x_i, y_j),
53 \\
54 &
55 H(x, y) = \begin{cases} 1 & (x < y) \\ 1/2 & (x = y) \end{cases}
56 \\
57 &
58 H^x_i = \sum_{j=1}^n H(y_j, x_i), \quad
59 H^y_j = \sum_{i=1}^m H(x_i, y_j),
60 \\
61 &
62 \bar{H}^x = \frac{1}{m} \sum_{i=1}^m H^x_i = n - n\hat{p},
63 \\
64 &
65 \bar{H}^y = \frac{1}{n} \sum_{j=1}^n H^y_j = m\hat{p}.
66 \end{aligned}
67 ````

68 この函数は以下達の named tuple で返す:
69
70 ````math
71 \begin{aligned}
72 &
73 \mathbf{phat} =
74 \hat{p} = \frac{\bar{H}^x - \bar{H}^y}{m + n},
75 \\
76 &
77 \mathbf{sx2} =
78 \hat{\sigma}_x^2 = \frac{1}{n^2} \frac{1}{m-1} \sum_{i=1}^m (H^x_i - \bar{H}^x)^2,
79 \\
80 &
81 \mathbf{sy2} =
82 \hat{\sigma}_y^2 = \frac{1}{m^2} \frac{1}{n-1} \sum_{j=1}^n (H^y_j - \bar{H}^y)^2,
83 \end{aligned}
84 ````
```

```

78  \\ &
79  \mathrm{sehat} =
80  \widehat{\mathrm{se}} = \sqrt{\frac{\hat{\sigma}_x^2}{m} + \frac{\hat{\sigma}_y^2}{n}},
81  \\ &
82  \mathrm{tvalue} = t = \frac{\hat{p} - p}{\widehat{\mathrm{se}}},
83  \\ &
84  \mathrm{df} =
85  \nu =
86  \frac{\left(\hat{\sigma}_x^2/m + \hat{\sigma}_y^2/n\right)^2}{\frac{\left(\hat{\sigma}_x^2/m\right)^2}{m-1} +
87  \frac{\left(\hat{\sigma}_y^2/n\right)^2}{n-1}},
88  },
89  \\ &
90  \mathrm{pvalue} =
91  2\mathit{ccdf}(\mathit{TDist}(\nu), |t|).
92  \end{aligned}
93  """
94  function statistics_brunner_munzel(X, Y,
95      Hx = similar(X, Float64),
96      Hy = similar(Y, Float64);
97      p = 1/2
98      )
99      m, n = length(X), length(Y)
100     for (i, x) in pairs(X)
101         Hx[i] = sum(h_brunner_munzel(y, x) for y in Y)
102     end
103     for (j, y) in pairs(Y)
104         Hy[j] = sum(h_brunner_munzel(x, y) for x in X)
105     end
106     phat = (mean(Hy) - mean(Hx) + n)/(m + n)
107     sx2, sy2 = var(Hx)/n^2, var(Hy)/m^2
108     sehat = \sqrt(sx2/m + sy2/n)
109     tvalue = (phat - p)/sehat
110     df = safediv((sx2/m + sy2/n)^2, (sx2/m)^2/(m-1) + (sy2/n)^2/(n-1))
111     pvalue = (df != 0 && isfinite(df)) ? 2ccdf(TDist(df), abs(tvalue)) : zero(df)
112     (; phat, sx2, sy2, sehat, tvalue, df, pvalue)
113  end
114
115  @doc raw"""
116      pvalue_brunner_munzel(X, Y,
117          Hx = similar(X, Float64),
118          Hy = similar(Y, Float64);
119          p = 1/2
120      )
121
122  この函数はBrunner-Munzel検定のP値 `pvalue` を返す.
123  """
124
125  function pvalue_brunner_munzel(X, Y,
126      Hx = similar(X, Float64),
127      Hy = similar(Y, Float64);
128      p = 1/2
129      )
130      statistics_brunner_munzel(X, Y, Hx, Hy; p).pvalue
131  end
132
133  """
134      tieshift(X::AbstractVector, Y::AbstractVector; p = 1/2)
135
136  この函数は `phat_brunner_munzel(X, Y .+ a)` の値が `p` に等しくなる `a` を返す.
137  """
138  function tieshift(X::AbstractVector, Y::AbstractVector; p = 1/2)
139      shiftmin = minimum(X) - maximum(Y) - 0.1
140      shiftmax = maximum(X) - minimum(Y) + 0.1
141      find_zero(a → phat_brunner_munzel(X, Y .+ a) - p, (shiftmin, shiftmax))
142  end
143
144  @doc raw"""
145      brunner_munzel(X, Y,
146          Hx = similar(X, Float64),
147          Hy = similar(Y, Float64),
148          Ytmp = similar(Y, Float64);
149          p = 1/2,
150          α = 0.05,
151          maxsplit = 30
152      )
153
154  """

```

```

156
157 この函数はBrunner-Munzel検定を実行する。 詳細は以下の通り。
158
159 この函数は `phat`, `sehat`, `tvalue`, `df`, `p`, `pvalue`, `α` および \
160 以下達の named tuple を返す。
161
162 ```\math
163 \begin{aligned}
164 & \\
165 \mathrm{confint\_p} = (\text{$p$ の信頼度 $1-\alpha$ の信頼区間}), \\
166 & \\
167 \mathrm{confint\_shift} = (\text{2つの集団が互角になるようなシフトの信頼度 $1-\alpha$ の信頼区間}), \\
168 & \\
169 \mathrm{pvalue\_shift} = ($\mathrm{confint\_shift}$ の計算で使われたP値函数), \\
170 & \\
171 \mathrm{shiftthat} = (\text{2つの集団が互角になるようなシフトの点推定値}). \\
172 \end{aligned}
173
174
175 さらに, $\mathrm{shiftmin}$, $\mathrm{shiftmax}$ はデータから推定されるシフトの下限と上限。
176
177 """
178 function brunner_munzel(X, Y,
179     Hx = similar(X, Float64),
180     Hy = similar(Y, Float64),
181     Ytmp = similar(Y, Float64);
182     p = 1/2,
183     α = 0.05,
184     maxsplit = 30
185 )
186     (; phat, sehat, tvalue, df, pvalue) = statistics_brunner_munzel(X, Y, Hx, Hy; p)
187
188     c = df == 0 ? Inf : quantile(TDist(df), 1 - α/2)
189     confint_p = [max(0, phat - c*sehat), min(1, phat + c*sehat)]
190
191     function pvalue_shift(a)
192         @. Ytmp = Y + a
193         pvalue_brunner_munzel(X, Ytmp, Hx, Hy; p)
194     end
195     shiftmin = minimum(X) - maximum(Y) - 0.1
196     shiftmax = maximum(X) - minimum(Y) + 0.1
197     shifthat = tieshift(X, Y; p)
198     confint_shift = [
199         find_zero(a → pvalue_shift(a) - α, (shiftmin, shifthat))
200         find_zero(a → pvalue_shift(a) - α, (shifthat, shiftmax))
201     ]
202
203     (; phat, sehat, tvalue, df, p, pvalue, α, confint_p,
204         confint_shift, pvalue_shift, shifthat, shiftmin, shiftmax)
205 end
206
207 function show_plot_brunner_munzel(X, Y,
208     Hx = similar(X, Float64),
209     Hy = similar(Y, Float64),
210     Ytmp = similar(Y, Float64);
211     p = 1/2,
212     α = 0.05,
213     showXY = false,
214     kwargs...
215 )
216     showXY && (@show X Y)
217     (; phat, sehat, tvalue, df, p, pvalue, α, confint_p,
218         confint_shift, pvalue_shift, shifthat, shiftmin, shiftmax) =
219         brunner_munzel(X, Y, Hx, Hy, Ytmp; p, α)
220     pprint(; phat, sehat, tvalue, df, p, pvalue, α, confint_p,
221         confint_shift, shifthat))
222     println()
223     @show median(X) median(Y)
224     plot(pvalue_shift, shiftmin, shiftmax; label="")
225     vline!([tieshift(X, Y)]; label="", ls=:dash)
226     title!("P-value function of shift")
227     plot!(ytick=0:0.05:1)
228     plot!(; kwargs...)
229 end

```

Out[6]: `show_plot_brunner_munzel (generic function with 4 methods)`

```
In [7]: 1 @doc h_brunner_munzel
```

```
Out[7]: h_brunner_munzel(x, y)
```

この函数は, $x < y$ のとき 1.0 を, $x = y$ のとき 0.5 を返す.

```
In [8]: 1 @doc phat_brunner_munzel
```

```
Out[8]: phat_brunner_munzel(X, Y)
```

まず以下のようにおく:

$$H(x, y) = \begin{cases} 1 & (x < y) \\ 1/2 & (x = y), \end{cases}$$

$$m = \text{length}(X), \quad n = \text{length}(Y), \quad x_i = X[i], \quad y_j = Y[j]$$

この函数は次の \hat{p} を返す:

$$\hat{p} = \frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n H(x_i, y_j).$$

In [9]: 1 @doc statistics_brunner_munzel

Out[9]:

```
statistics_brunner_munzel(X, Y,
    Hx = similar(X, Float64),
    Hy = similar(Y, Float64);
    p = 1/2
)
```

この函数はデータ X , Y について, Brunner-Munzel検定関係の統計量達を計算する. 詳細は以下の通り.

函数 $H(x, y)$ と \hat{p} , H_i^x , H_j^y , \bar{H}^x , \bar{H}^y を次のように定める:

$$m = \text{length}(X), \quad n = \text{length}(Y), \quad x_i = X[i], \quad y_j = Y[j],$$

$$\hat{p} = \frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n H(x_i, y_j),$$

$$H(x, y) = \begin{cases} 1 & (x < y) \\ 1/2 & (x = y), \end{cases}$$

$$H_i^x = \sum_{j=1}^n H(y_j, x_i), \quad H_j^y = \sum_{i=1}^m H(x_i, y_j),$$

$$\bar{H}^x = \frac{1}{m} \sum_{i=1}^m H_i^x = n - n\hat{p},$$

$$\bar{H}^y = \frac{1}{n} \sum_{j=1}^n H_j^y = m\hat{p}.$$

この函数は以下達の named tuple で返す:

$$\text{phat} = \hat{p} = \frac{\bar{H}^x - \bar{H}^y + n}{m + n},$$

$$\text{sx2} = \hat{\sigma}_x^2 = \frac{1}{n^2} \frac{1}{m-1} \sum_{i=1}^m (H_i^x - \bar{H}^x)^2,$$

$$\text{sy2} = \hat{\sigma}_y^2 = \frac{1}{m^2} \frac{1}{n-1} \sum_{j=1}^n (H_j^y - \bar{H}^y)^2,$$

$$\text{sehat} = \widehat{\text{se}} = \sqrt{\frac{\hat{\sigma}_x^2}{m} + \frac{\hat{\sigma}_y^2}{n}},$$

$$\text{tvalue} = t = \frac{\hat{p} - p}{\widehat{\text{se}}},$$

$$\text{df} = v = \frac{\left(\hat{\sigma}_x^2/m + \hat{\sigma}_y^2/n\right)^2}{\frac{\left(\hat{\sigma}_x^2/m\right)^2}{m-1} + \frac{\left(\hat{\sigma}_y^2/n\right)^2}{n-1}},$$

$$\text{pvalue} = 2\text{ccdf}(\text{TDist}(v), |t|).$$

```
In [10]: 1 @doc brunner_munzel
```

```
Out[10]: brunner_munzel(X, Y,
    Hx = similar(X, Float64),
    Hy = similar(Y, Float64),
    Ytmp = similar(Y, Float64);
    p = 1/2,
    α = 0.05,
    maxsplit = 30
)
```

この函数はBrunner-Munzel検定を実行する。詳細は以下の通り。

この函数は `phat`, `sehat`, `tvalue`, `df`, `p`, `pvalue`, `α` および以下達の named tuple を返す。

`confint_p` = (p の信頼度 $1 - \alpha$ の信頼区間),
`confint_shift` = (2つの集団が互角になるようなシフトの信頼度 $1 - \alpha$ の信頼区間),
`pvalue_shift` = (\$`confint_shift$`の計算で使われた P 値函数),
`shifthat` = (2つの集団が互角になるようなシフトの点推定値)。

さらに, `shiftmin`, `shiftmax` はデータから推定されるシフトの下限と上限。

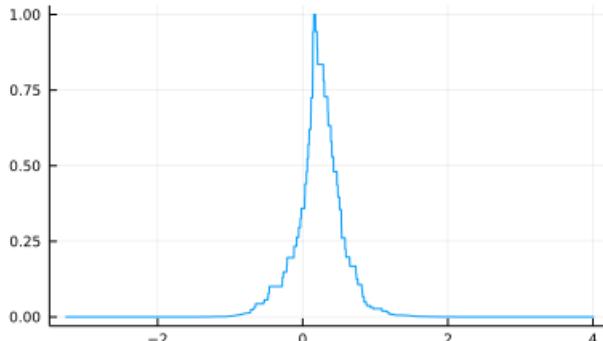
```
In [11]: 1 X = randn(10)
2 Y = randn(10)
3 @show shiftmin = minimum(X) - maximum(Y) - 1
4 @show shiftmax = maximum(X) - minimum(Y) + 1
5 pvalue_brunner_munzel(X, Y)
```

```
shiftmin = (minimum(X) - maximum(Y)) - 1 = -3.2716912875061364
shiftmax = (maximum(X) - minimum(Y)) + 1 = 4.017034981351518
```

```
Out[11]: 0.35885925554743014
```

```
In [12]: 1 plot(a → pvalue_brunner_munzel(X, Y .+ a), shiftmin, shiftmax;
2           label="", title="P-value function of shift")
```

```
Out[12]: P-value function of shift
```



2.2 よく使われているっぽいテストデータで正しく実装されているかを確認

<https://okumuralab.org/~okumura/stat/brunner-munzel.html> (<https://okumuralab.org/~okumura/stat/brunner-munzel.html>)

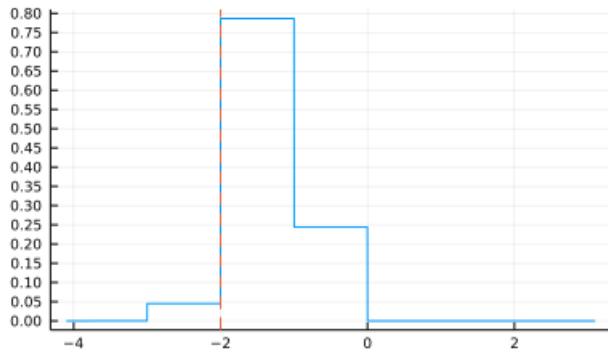
```
x = c(1,2,1,1,1,1,1,1,1,2,4,1,1)
y = c(3,3,4,3,1,2,3,1,1,5,4)
brunnermunzel.test(x, y)

data: x and y
Brunner-Munzel Test Statistic = 3.1375, df = 17.683, p-value = 0.005786
95 percent confidence interval:
 0.5952169 0.9827052
sample estimates:
P(X<Y)+.5*P(X=Y)
 0.788961
```

```
In [13]: 1 X = [1,2,1,1,1,1,1,1,1,1,2,4,1,1]
2 Y = [3,3,4,3,1,2,3,1,1,5,4]
3 show_plot_brunner_munzel(X, Y)
```

```
(phat = 0.788961038961039,
sehat = 0.09210009046816862,
tvalue = 3.1374674823029505,
df = 17.682841979481545,
p = 0.5,
pvalue = 0.005786208666151463,
alpha = 0.05,
confint_p = [0.5952168642537363, 0.9827052136683416],
confint_shift = [-2.0000000000000004, -5.551115123125783e-17],
shiftthat = -1.9999999999999998)
median(X) = 1.0
median(Y) = 3.0
```

Out[13]: P-value function of shift



```
In [14]: 1 X = [1,2,1,1,1,1,1,1,1,1,2,4,1,1]
2 Y = [3,3,4,3,1,2,3,1,1,5,4]
3 @rput X Y
4 R"""
5 library(lawstat)
6 brunner.munzel.test(X, Y)
7 """
```

Out[14]: RObject{VecSxp}

```
Brunner-Munzel Test

data: X and Y
Brunner-Munzel Test Statistic = 3.1375, df = 17.683, p-value = 0.005786
95 percent confidence interval:
0.5952169 0.9827052
sample estimates:
P(X<Y)+.5*p(X=Y)
0.788961
```

このように Brunner-Munzel 検定は R では [lawstat](https://cran.r-project.org/package=lawstat) (<https://cran.r-project.org/package=lawstat>) パッケージの [brunner.munzel.test](https://rdrr.io/cran/lawstat/man/brunner.munzel.test.html) (<https://rdrr.io/cran/lawstat/man/brunner.munzel.test.html>) で使える。

2.3 組み合わせの生成子

```
In [15]: 1 """
2     complementcomb!(complcomb::AbstractVector, comb::AbstractVector)
3
4 'comb' が {1,2,...,N} から重複無しに m 個を選ぶ組み合わせを表す配列であり, 'comb' の中で数は小さ
5 このとき, この函数は配列 'complcomb' に配列 'comb' の補集合を格納し, 'complcomb' を返す.
6
7 この函数はメモリ割り当てゼロで実行される.
8 """
9
10 function complementcomb!(complcomb::AbstractVector, comb::AbstractVector)
11     N = length(comb) + length(complcomb)
12     k = 0
13     a = 0
14     @inbounds for b in comb
15         for i in a+1:b-1
16             k += 1
17             complcomb[k] = i
18         end
19         a = b
20     end
21     @inbounds for i in a+1:N
22         k += 1
23         complcomb[k] = i
24     end
25     complcomb
26 end
27 """
28     complementcomb(N, comb::AbstractVector)
29
30 'comb' が {1,2,...,N} から重複無しに m 個を選ぶ組み合わせを表す配列であり, 'comb' の中で数は小さ
31 この函数は 'comb' の補集合の配列を返す.
32
33 この函数は返り値の配列の分だけのメモリ割り当てを行う.
34 """
35
36 complementcomb(N, comb::AbstractVector) =
37     complementcomb!(similar(comb, N - length(comb)), comb)
```

Out[15]: complementcomb

In [16]: 1 @doc complementcomb!

Out[16]: complementcomb!(complcomb::AbstractVector, comb::AbstractVector)

comb が {1,2,...,N} から重複無しに m 個を選ぶ組み合わせを表す配列であり, comb の中で数は小さな順に並んでいるとし, complcomb は長さ N - m の配列であると仮定する.

このとき, この函数は配列 complcomb に配列 comb の補集合を格納し, complcomb を返す.

この函数はメモリ割り当てゼロで実行される.

In [17]: 1 @doc complementcomb

Out[17]: complementcomb(N, comb::AbstractVector)

comb が {1,2,...,N} から重複無しに m 個を選ぶ組み合わせを表す配列であり, comb の中で数は小さな順に並んでいると仮定する.

この函数は comb の補集合の配列を返す.

この函数は返り値の配列の分だけのメモリ割り当てを行う.

In [18]: 1 N = 10
2 comb = [2, 4, 5, 8]
3 ccomb = similar(comb, N - length(comb))
4 @btime complementcomb!(\$ccomb, \$comb);

9.810 ns (0 allocations: 0 bytes)

```
In [19]: 1 N, m = 5, 3
          2 ccomb = Vector{Int}(undef, N-m)
          3 [(copy(comb), copy(complementcomb!(ccomb, comb))) for comb in mycombinations(1:N, m)]
```

Out[19]: 10-element Vector{Tuple{Vector{Int64}, Vector{Int64}}}:

```
([1, 2, 3], [4, 5])
([1, 2, 4], [3, 5])
([1, 2, 5], [3, 4])
([1, 3, 4], [2, 5])
([1, 3, 5], [2, 4])
([1, 4, 5], [2, 3])
([2, 3, 4], [1, 5])
([2, 3, 5], [1, 4])
([2, 4, 5], [1, 3])
([3, 4, 5], [1, 2])
```

```
In [20]: 1 N, m = 5, 3
          2 ccomb = Vector{Int}(undef, N-m)
          3 [(copy(comb), complementcomb(N, comb)) for comb in mycombinations(1:N, m)]
```

Out[20]: 10-element Vector{Tuple{Vector{Int64}, Vector{Int64}}}:

```
([1, 2, 3], [4, 5])
([1, 2, 4], [3, 5])
([1, 2, 5], [3, 4])
([1, 3, 4], [2, 5])
([1, 3, 5], [2, 4])
([1, 4, 5], [2, 3])
([2, 3, 4], [1, 5])
([2, 3, 5], [1, 4])
([2, 4, 5], [1, 3])
([3, 4, 5], [1, 2])
```

2.4 Brunner-Munzel検定のPermutation版の実装

In [21]:

```
1 """
2     permutation_tvalues_brunner_munzel(X, Y,
3         XandY = Vector{Float64}(undef, length(X)+length(Y)),
4         Tval = Vector{Float64}(undef, binomial(length(X)+length(Y), length(X))),
5         Hx = similar(X, Float64),
6         Hy = similar(Y, Float64)
7     )
8
9 Brunner-Munzel検定のt値を '[X; Y]' から \
10 インデックスの重複無しに 'length(X)' 個取る組み合わせと \
11 その補集合への分割のすべてについて計算して, 'Tval' に格納して返す.
12 """
13 function permutation_tvalues_brunner_munzel(X, Y,
14     XandY = Vector{Float64}(undef, length(X)+length(Y)),
15     Tval = Vector{Float64}(undef, binomial(length(X)+length(Y), length(X))),
16     Hx = similar(X, Float64),
17     Hy = similar(Y, Float64),
18     ccomb = Vector{Int}(undef, length(Y))
19 )
20 m, n = length(X), length(Y)
21 N = m + n
22 @views XandY[1:m] .= X
23 @views XandY[m+1:N] .= Y
24 for (k, comb) in enumerate(mycombinations(1:N, m))
25     complementcomb!(ccomb, comb)
26     Tval[k] = statistics_brunner_munzel(
27         view(XandY, comb), view(XandY, ccomb), Hx, Hy).tvalue
28 end
29 Tval
30 end
31 """
32 pvalue_brunner_munzel_perm(X, Y,
33     Tval = permutation_tvalues_brunner_munzel(X, Y),
34     tval = statistics_brunner_munzel(X, Y).tvalue;
35     le = ≈
36 )
37
38 Brunner-Munzel検定のpermutation版のP値を返す.
39 """
40 function pvalue_brunner_munzel_perm(X, Y,
41     Tval = permutation_tvalues_brunner_munzel(X, Y),
42     tval = statistics_brunner_munzel(X, Y).tvalue;
43     le = ≈
44 )
45     pvalue_perm = mean(T → le(abs(tval), abs(T)), Tval)
46 end
47 end
```

Out[21]: pvalue_brunner_munzel_perm

In [22]:

```
1 @doc permutation_tvalues_brunner_munzel
```

Out[22]:

```
permutation_tvalues_brunner_munzel(X, Y,
    XandY = Vector{Float64}(undef, length(X)+length(Y)),
    Tval = Vector{Float64}(undef, binomial(length(X)+length(Y), length(X))),
    Hx = similar(X, Float64),
    Hy = similar(Y, Float64)
)
```

Brunner-Munzel検定のt値を [X; Y] からインデックスの重複無しに length(X) 個取る組み合わせとその補集合への分割のすべてについて計算して, Tval に格納して返す.

In [23]:

```
1 @doc pvalue_brunner_munzel_perm
```

Out[23]:

```
pvalue_brunner_munzel_perm(X, Y,
    Tval = permutation_tvalues_brunner_munzel(X, Y),
    tval = statistics_brunner_munzel(X, Y).tvalue;
    le = ≈
)
```

Brunner-Munzel検定のpermutation版のP値を返す.

```

bm = brunner.munzel.test(x, y)$statistic
n1 = length(x)
n2 = length(y)
N = n1 + n2
xandy = c(x, y)
foo = function(X) {
  brunner.munzel.test(xandy[X], xandy[-X])$statistic
}
z = combn(1:N, n1, foo)
mean(abs(z) >= abs(bm))

```

結果は 0.008037645 となりました。

In [24]:

```

1 X = [1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 2, 4, 1, 1]
2 Y = [3, 3, 4, 3, 1, 2, 3, 1, 1, 5, 4]
3 @show X Y
4 @show m, n = length(X), length(Y)
5
6 Tval = @time permutation_tvalues_brunner_munzel(X, Y)
7 @show pvalue_brunner_munzel_perm(X, Y, Tval)
8 stephist(Tval; norm=true, bin=101, label="", title="permutation t-values")

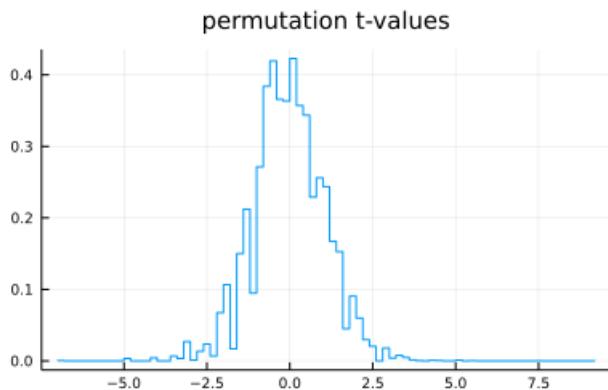
```

```

X = [1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 2, 4, 1, 1]
Y = [3, 3, 4, 3, 1, 2, 3, 1, 1, 5, 4]
(m, n) = (length(X), length(Y)) = (14, 11)
4.103247 seconds (1.01 M allocations: 84.027 MiB, 0.41% gc time, 7.10% compilation time)
pvalue_brunner_munzel_perm(X, Y, Tval) = 0.008037645264055279

```

Out[24]:



2.5 permutation版が正しく実装されているかの確認

- <https://github.com/toshi-ara/brunnermunzel/issues/14> (<https://github.com/toshi-ara/brunnermunzel/issues/14>)
- <https://github.com/toshi-ara/brunnermunzel/files/4395032/mwe.R.zip> (<https://github.com/toshi-ara/brunnermunzel/files/4395032/mwe.R.zip>)

追記 2022-08-06: <https://twitter.com/TA25140989/status/1555825941451923457>

(<https://twitter.com/TA25140989/status/1555825941451923457>) を参照せよ。

<https://github.com/toshi-ara/brunnermunzel/tree/development> (<https://github.com/toshi-ara/brunnermunzel/tree/development>) の修正版の brunnermunzel パッケージをインストールし直した。以下のセルの実行結果が変わるはずなので、その記録を残しておく。

以下の記録を見なくても、

- <https://github.com/genkuroki/public/blob/e4faafc52721b63876b3b705f9450eade3c902f5/0034/Brunner-Munzel.ipynb> (<https://github.com/genkuroki/public/blob/e4faafc52721b63876b3b705f9450eade3c902f5/0034/Brunner-Munzel.ipynb>)

で閲覧できるが、わざわざ見に行くのも面倒なのでこのファイルにも記録を残しておく。

以前の実行結果:

```
@show pval_J - pval_J_le;
```

```
pval_J - pval_J_le = [0.0, 0.0, 0.007936507936507936, 0.023809523809523808, 0.023809523809  
523808, 0.0793650793650793, 0.0, 0.0, 0.007936507936507936, 0.0357142857142857, 0.0, 0.015  
873015873015928, 0.015873015873015872, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0, 0.0, 0.0079365  
07936507936, 0.0, 0.023809523809523836, 0.015873015873015928, 0.011904761904761918, 0.0,  
0.015873015873015928, 0.0, 0.003968253968253968, 0.0, 0.0, 0.0, 0.0, 0.00793650793650793  
6, 0.0, 0.015873015873015928, 0.0, 0.015873015873015928, 0.0, 0.0, 0.0, 0.0079365079365079  
08, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.0, 0.007936507936507936,  
0.007936507936507936, 0.011904761904761918, 0.007936507936507936, 0.03968253968253965, 0.  
0, 0.007936507936507936, 0.0, 0.05555555555555547, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.015873015873015872, 0.015873015873015872, 0.0, 0.015873015873015928,  
0.0, 0.0, 0.0, 0.0, 0.0, 0.015873015873015928, 0.007936507936507964, 0.00793650793650  
7964, 0.0, 0.003968253968253954, 0.0, 0.0, 0.0, 0.015873015873015872, 0.0, 0.0, 0.03571428  
57142857, 0.015873015873015872, 0.0, 0.023809523809523725, 0.0, 0.023809523809523808, 0.0,  
0.0, 0.0, 0.0, 0.0]
```

```
idx = @show findall(pval_J .!= pval_J_le)  
length(idx)
```

```
findall(pval_J .!= pval_J_le) = [3, 4, 5, 6, 9, 10, 12, 13, 15, 19, 21, 22, 23, 25, 27, 3  
2, 34, 36, 40, 42, 44, 47, 48, 49, 50, 51, 53, 55, 68, 69, 71, 78, 79, 80, 82, 86, 89, 90,  
92, 94]
```

40

```
@show pval_R - pval_J_le;
```

```
pval_R - pval_J_le = [0.0, 0.0, 0.007936507936507936, 0.023809523809523808, 0.023809523809  
523808, 0.0793650793650793, 0.0, 0.0, 0.007936507936507936, 0.0357142857142857, 0.0, 0.015  
873015873015928, 0.015873015873015872, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0, 0.0, 0.0079365  
07936507936, 0.0, 0.023809523809523836, 0.007936507936507908, 0.011904761904761918, 0.0,  
0.015873015873015928, 0.0, 0.003968253968253968, 0.0, 0.0, 0.0, 0.0, 0.00793650793650793  
6, 0.0, 0.00793650793650802, 0.0, 0.015873015873015928, 0.0, 0.0, 0.0, 0.00793650793650790  
8, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.0, 0.007936507936507936,  
0.007936507936507936, 0.011904761904761918, 0.007936507936507936, 0.03968253968253965, 0.  
0, 0.007936507936507936, 0.0, 0.05555555555555547, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.015873015873015872, 0.015873015873015872, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.007936507936507908, 0.007936507936507964, 0.007936507936507964, 0.0,  
0.03968253968253954, 0.0, 0.0, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0357142857142857, 0.01  
5873015873015872, 0.0, 0.023809523809523725, 0.0, 0.023809523809523808, 0.0, 0.0, 0.0, 0.  
0, 0.0, 0.0]
```

```
idx = @show findall(pval_R .!= pval_J_le)  
length(idx)
```

```
findall(pval_R .!= pval_J_le) = [3, 4, 5, 6, 9, 10, 12, 13, 15, 19, 21, 22, 23, 25, 27, 3  
2, 34, 36, 40, 42, 44, 47, 48, 49, 50, 51, 53, 55, 68, 69, 78, 79, 80, 82, 86, 89, 90, 92,  
94]
```

39

```
@show pval_J - pval_R;
```

```
pval_J - pval_R = [0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.00793650793650802, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0, 0.0, 0.0, 0.0, 0.0, 0.007936507936507908, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.00793650793650802, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,  
0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0,
```

```
idx = @show findall(.!(pval_J .≈ pval_R))  
length(idx)
```

```
findall(.!(pval_J .≈ pval_R)) = [22, 34, 71, 78]
```

4

```
all(pval_J .≥ pval_R .≥ pval_J_le)
```

```
true
```

以前のコメントは以下の通り:

なるほど！

<https://github.com/toshi-ara/brunnermunzel/issues/14>

に書いてあるように，22, 34, 71 and 78 の4つで，値が一致していない。

○○以下または○○以上の判定を \$x \approx y\$ のときも true にする必要があるのだが，その部分で違いが生じているものと思われる。

現時点では <https://CRAN.R-project.org/package=brunnermunzel> にアクセスすると，

```
>Package ‘brunnermunzel’ was removed from the CRAN repository.  
>  
>Formerly available versions can be obtained from the [archive](https://cran.r-project.org/src/contrib/Archive/brunnermunzel/).  
>  
>Archived on 2022-03-04 as check problems were not corrected in time. , LENGTH_1 checks.  
>  
>A summary of the most recent check results can be obtained from the [check results archive](https://cran-archive.r-project.org/web/checks/2022/2022-03-04\_check\_results\_brunnermunzel.html).  
>  
>Please use the canonical form https://CRAN.R-project.org/package=brunnermunzel to link to  
this page.
```

と表示される。

In [25]:

```
1 R"""  
2 library(brunnermunzel)  
3 set.seed(1290)  
4 reps = 100  
5 xx = c()  
6 yy = c()  
7 pval_R = numeric(reps)  
8 for (i in seq_len(reps)){  
9   x = rnorm(5)  
10  y = rnorm(5)  
11  
12  xx = c(xx, x)  
13  yy = c(yy, y)  
14  
15  res_bm_perm ← brunnermunzel.permutation.test(x,y)  
16  pval_R[i] ← res_bm_perm$p.value  
17 }  
18 """  
19  
20 @rget xx yy pval_R  
21 XX = reshape(xx, 5, 100)  
22 YY = reshape(yy, 5, 100)  
23  
24 pval_J = zeros(100)  
25 pval_J_le = zeros(100)  
26 for i in 1:100  
27   pval_J[i] = pvalue_brunner_munzel_perm(XX[ :,i], YY[ :,i]; le = ≈)  
28   pval_J_le[i] = pvalue_brunner_munzel_perm(XX[ :,i], YY[ :,i]; le = ≤)  
29 end
```

```
In [26]: 1 @show pval_J - pval_J_le;
```

```
pval_J - pval_J_le = [0.0, 0.0, 0.007936507936507936, 0.023809523809523808, 0.023809523809523808, 0.0793650793650793, 0.0, 0.0, 0.007936507936507936, 0.0357142857142857, 0.0, 0.015873015873015928, 0.015873015873015872, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0, 0.007936507936507936, 0.0, 0.023809523809523836, 0.015873015873015928, 0.011904761904761918, 0.0, 0.015873015873015928, 0.0, 0.003968253968253968, 0.0, 0.0, 0.0, 0.0, 0.007936507936507936, 0.0, 0.015873015873015928, 0.0, 0.015873015873015928, 0.0, 0.0, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.0, 0.007936507936507936, 0.007936507936507936, 0.011904761904761918, 0.007936507936507936, 0.03968253968253965, 0.0, 0.007936507936507936, 0.0, 0.05555555555555547, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.015873015873015872, 0.015873015873015872, 0.0, 0.015873015873015928, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.015873015873015928, 0.007936507964, 0.007936507936507964, 0.0, 0.003968253968253954, 0.0, 0.0, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0, 0.0357142857142857, 0.015873015873015872, 0.0, 0.023809523809523725, 0.0, 0.023809523808, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0]
```

```
In [27]: 1 idx = @show findall(pval_J .!= pval_J_le)
2 length(idx)
```

```
findall(pval_J .!= pval_J_le) = [3, 4, 5, 6, 9, 10, 12, 13, 15, 19, 21, 22, 23, 25, 27, 32, 34, 36, 40, 42, 44, 47, 48, 49, 50, 51, 53, 55, 68, 69, 71, 78, 79, 80, 82, 86, 89, 90, 92, 94]
```

Out[27]: 40

```
In [28]: 1 @show pval_R - pval_J_le;
```

```
pval_R - pval_J_le = [0.0, 0.0, 0.007936507936507936, 0.023809523809523808, 0.023809523809523808, 0.0793650793650793, 0.0, 0.0, 0.007936507936507936, 0.0357142857142857, 0.0, 0.015873015873015928, 0.015873015873015872, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0, 0.007936507936507936, 0.0, 0.023809523809523836, 0.015873015873015928, 0.011904761904761918, 0.0, 0.015873015873015928, 0.0, 0.003968253968253968, 0.0, 0.0, 0.0, 0.0, 0.007936507936507936, 0.0, 0.015873015873015928, 0.0, 0.015873015873015928, 0.0, 0.0, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.007936507936507908, 0.0, 0.0, 0.007936507936507936, 0.007936507936507936, 0.011904761904761918, 0.007936507936507936, 0.03968253968253965, 0.0, 0.007936507936507936, 0.0, 0.05555555555555547, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.015873015873015872, 0.015873015873015872, 0.0, 0.015873015873015928, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.015873015873015928, 0.007936507964, 0.007936507936507964, 0.0, 0.003968253968253954, 0.0, 0.0, 0.0, 0.015873015873015872, 0.0, 0.0, 0.0357142857142857, 0.015873015873015872, 0.0, 0.023809523809523725, 0.0, 0.023809523808, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0]
```

```
In [29]: 1 idx = @show findall(pval_R .!= pval_J_le)
2 length(idx)
```

```
findall(pval_R .!= pval_J_le) = [3, 4, 5, 6, 9, 10, 12, 13, 15, 19, 21, 22, 23, 25, 27, 32, 34, 36, 40, 42, 44, 47, 48, 49, 50, 51, 53, 55, 68, 69, 71, 78, 79, 80, 82, 86, 89, 90, 92, 94]
```

Out[29]: 40

```
In [30]: 1 @show pval_J - pval_R;
```

```
In [31]: 1 idx = @show findall(.!{pval_J ≈ pval_R})
2 length(idx)
```

```
findall(.!(pval_J .≈ pval_R)) = Int64[]
```

Out[31]: 0

```
In [32]: 1 all(pval_J .>= pval_R .>= pval_J_le)
```

Out[32]: true

2022-08-06: やった! 値が完全に一致した! permutation版Brunner-Munzel検定について、

- <https://github.com/toshi-ara/brunnermunzel/tree/development> (<https://github.com/toshi-ara/brunnermunzel/tree/development>)

の実装と私による実装の計算結果は以下の場合において完全に一致している。

3 計算例

In [33]:

```
1 m, n = 10, 10
2 X, Y = rand(Normal(0, 1), m), rand(Normal(0, 2), n)
3 @show pval_brmu = pvalue_brunner_munzel(X, Y)
4 @show pval_perm = pvalue_brunner_munzel_perm(X, Y);

pval_brmu = pvalue_brunner_munzel(X, Y) = 0.37377292369136617
pval_perm = pvalue_brunner_munzel_perm(X, Y) = 0.35041893091428694
```

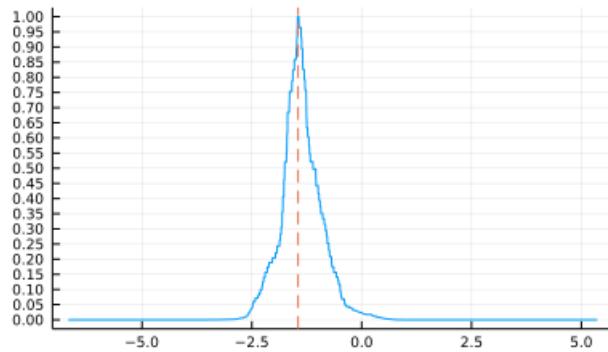
In [34]:

```
1 Random.seed!(4)
2
3 m, n = 10, 20
4 X, Y = rand(Normal(0, 1), m), rand(Normal(0, 2), n)
5 show_plot_brunner_munzel(X, Y)
```

```
(phat = 0.75,
sehat = 0.10056855914111178,
tvalue = 2.485866379463735,
df = 18.385702920172452,
p = 0.5,
pvalue = 0.022742714988590425,
α = 0.05,
confint_p = [0.5390305665719614, 0.9609694334280386],
confint_shift = [-2.4770370893730127, -0.3717172074086195],
shiftthat = -1.4469029166494445)
median(X) = -0.5058729691113518
median(Y) = 1.1395412005826189
```

Out[34]:

P-value function of shift



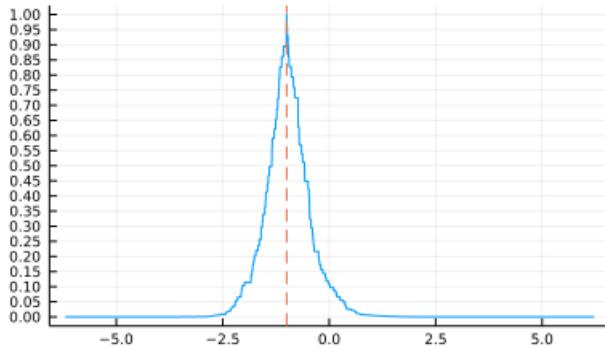
In [35]:

```
1 m, n = 10, 20
2 X, Y = rand(Normal(0, 1), m), rand(Normal(0, 2), n)
3 show_plot_brunner_munzel(X, Y)

(phat = 0.675,
 sehat = 0.1049296617650541,
 tvalue = 1.667783895004246,
 df = 27.705176944214784,
 p = 0.5,
 pvalue = 0.10662428006873317,
 alpha = 0.05,
 confint_p = [0.45995823964489857, 0.8900417603551015],
 confint_shift = [-2.2060635904773678, 0.3296238740154929],
 shifthat = -0.999368724114565)
median(X) = -0.17409123783026362
median(Y) = 0.8382134149102431
```

Out[35]:

P-value function of shift



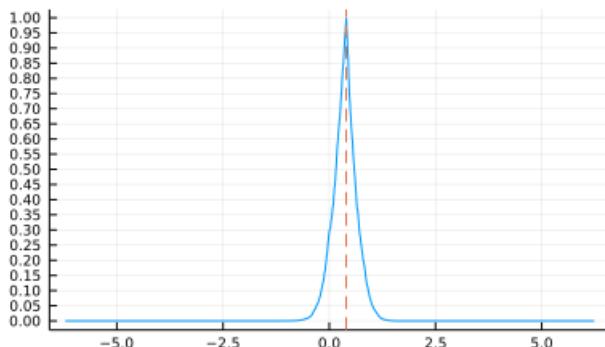
In [36]:

```
1 m, n = 100, 50
2 X, Y = rand(Normal(0, 1), m), rand(Normal(0, 2), n)
3 show_plot_brunner_munzel(X, Y)

(phat = 0.4397999999999997,
 sehat = 0.05714122667226539,
 tvalue = -1.0535300606211744,
 df = 60.575330843208235,
 p = 0.5,
 pvalue = 0.2962824213806768,
 alpha = 0.05,
 confint_p = [0.3255228528192483, 0.5540771471807516],
 confint_shift = [-0.3011419367780337, 1.0179208275627627],
 shifthat = 0.3967840840930079)
median(X) = -0.0536449473157313
median(Y) = -0.6163649344394873
```

Out[36]:

P-value function of shift



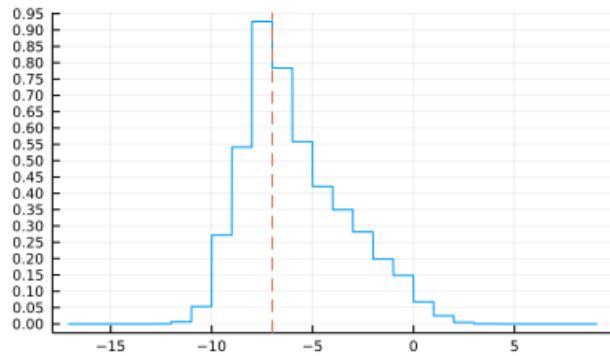
In [37]:

```
1 Random.seed!(4)
2
3 m, n = 10, 20
4 X, Y = rand(1:m, m), rand(1:n, n)
5 show_plot_brunner_munzel(X, Y)
```

```
(phat = 0.6725,
sehat = 0.1005725567999867,
tvalue = 1.715179622439735,
df = 20.71835365424782,
p = 0.5,
pvalue = 0.10123190720539166,
α = 0.05,
confint_p = [0.46317465971191163, 0.8818253402880883],
confint_shift = [-10.999999999999998, 0.9999999999999997],
shiftthat = -7.000000000000001)
median(X) = 6.5
median(Y) = 14.5
```

Out[37]:

P-value function of shift



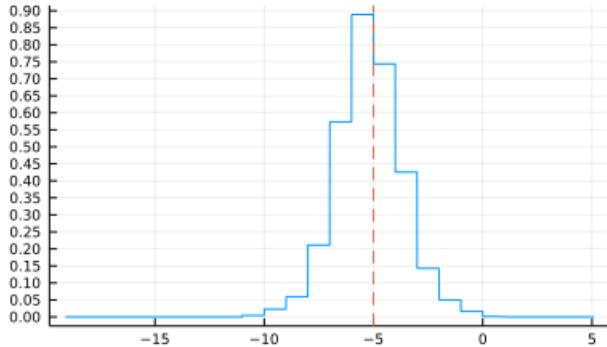
In [38]:

```
1 m, n = 10, 20
2 X, Y = rand(1:m, m), rand(1:n, n)
3 show_plot_brunner_munzel(X, Y)

(phat = 0.765,
 sehat = 0.08626255311649104,
 tvalue = 3.0720166564295583,
 df = 27.60349766038036,
 p = 0.5,
 pvalue = 0.004741546491682432,
 alpha = 0.05,
 confint_p = [0.5881847509519754, 0.9418152490480246],
 confint_shift = [-8.999999999999998, -1.999999999999993],
 shifthat = -5.0)
median(X) = 5.5
median(Y) = 12.0
```

Out[38]:

P-value function of shift



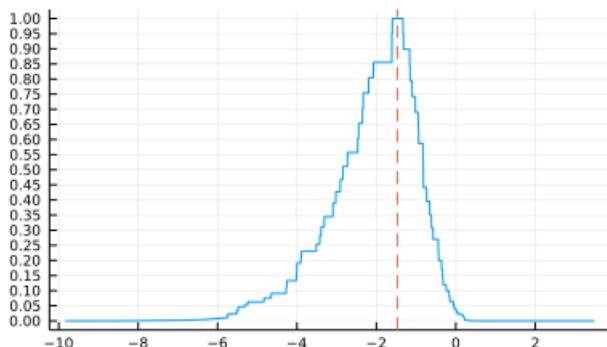
In [39]:

```
1 distx, disty = LogNormal(), LogNormal(1)
2 m, n = 10, 10
3 X, Y = rand(distx, m), rand(disty, n)
4 show_plot_brunner_munzel(X, Y)

(phat = 0.75,
 sehat = 0.11205157542647741,
 tvalue = 2.231115439907736,
 df = 18.0,
 p = 0.5,
 pvalue = 0.03863103202434314,
 alpha = 0.05,
 confint_p = [0.5145883755427825, 0.9854116244572175],
 confint_shift = [-5.318179432610124, -0.05841508479073754],
 shifthat = -1.4733410643825495)
median(X) = 1.3089930416807416
median(Y) = 2.9790059563220423
```

Out[39]:

P-value function of shift

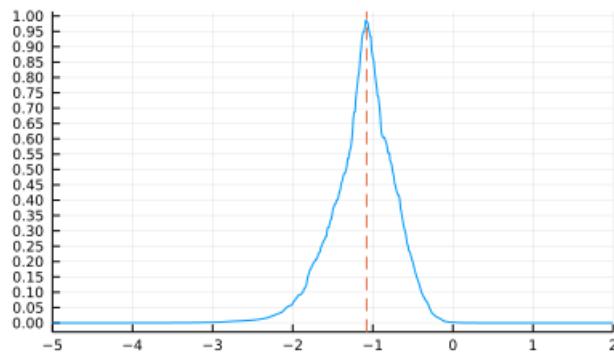


```
In [40]: 1 distx, disty = LogNormal(), LogNormal(1)
2 m, n = 40, 40
3 X, Y = rand(distx, m), rand(disty, n)
4 show_plot_brunner_munzel(X, Y; xlim=(-5, 2))
```

```
(phat = 0.6912499999999999,
sehat = 0.05960002903974512,
tvalue = 3.208891053936604,
df = 75.84080882915966,
p = 0.5,
pvalue = 0.001953574321225763,
α = 0.05,
confint_p = [0.5725422250913458, 0.8099577749086541],
confint_shift = [-2.07997163873439, -0.29464656357396524],
shiftthat = -1.0768936945378669)
median(X) = 1.044779302249417
median(Y) = 2.2527593566191237
```

Out[40]:

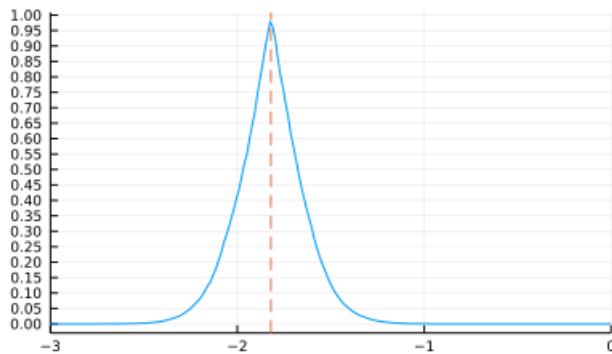
P-value function of shift



```
In [41]: 1 distx, disty = LogNormal(), LogNormal(1)
2 m, n = 160, 160
3 X, Y = rand(distx, m), rand(disty, n)
4 show_plot_brunner_munzel(X, Y; xlim=(-3, 0))
```

```
(phat = 0.79234375,
sehat = 0.025077139980214335,
tvalue = 11.657778767062629,
df = 315.6587283148897,
p = 0.5,
pvalue = 2.3287816342563994e-26,
α = 0.05,
confint_p = [0.7430042840809774, 0.8416832159190226],
confint_shift = [-2.250313266911184, -1.404197630202607],
shiftthat = -1.8203983580938916)
median(X) = 0.8417596982690712
median(Y) = 2.974128627007357
```

Out[41]: P-value function of shift



4 Brunner-Munzel検定とWelchのt検定の比較

4.1 第一種の過誤の確率

In [42]:

```

1 function sim_brunner_mumzel();
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10,
3     L = 10^6)
4     pval_bm = Vector{Float64}(undef, L)
5     tmpX = [Vector{Float64}(undef, m) for _ in 1:nthreads()]
6     tmpY = [Vector{Float64}(undef, n) for _ in 1:nthreads()]
7     tmpHx = [Vector{Float64}(undef, m) for _ in 1:nthreads()]
8     tmpHy = [Vector{Float64}(undef, n) for _ in 1:nthreads()]
9     @threads for i in 1:L
10         X = rand!(distx, tmpX[threadid()])
11         Y = rand!(disty, tmpY[threadid()])
12         pval_bm[i] = pvalue_brunner_munzel(X, Y, tmpHx[threadid()], tmpHy[threadid()])
13     end
14     ecdf(pval_bm)
15 end
16
17 function sim_welch();
18     distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10,
19     L = 10^6)
20     pval_w = Vector{Float64}(undef, L)
21     tmpX = [Vector{Float64}(undef, m) for _ in 1:nthreads()]
22     tmpY = [Vector{Float64}(undef, n) for _ in 1:nthreads()]
23     @threads for i in 1:L
24         X = rand!(distx, tmpX[threadid()])
25         Y = rand!(disty, tmpY[threadid()])
26         pval_w[i] = pvalue_welch(X, Y)
27     end
28     ecdf(pval_w)
29 end
30
31 function printcompact(io, xs...)
32     print(IOContext(io, :compact => true), xs...)
33 end
34
35 function distname(dist)
36     replace(sprint(printcompact, dist), r"\{\^\}\*\\"=>"")
37 end
38
39 function plot_ecdf(ecdf_pval, distx, disty, m, n, a;
40     testname = "", kwargs...)
41     plot(p → ecdf_pval(p), 0, 0.1; label="ecdf of P-values")
42     plot!([0, 0.1], [0, 0.1]; label="", ls=:dot, c=:black)
43     plot!(legend=:topleft)
44     plot!(xtick=0:0.01:0.1, ytick=0:0.01:1)
45     plot!(xguide="nominal significance level  $\alpha$ ",
46           yguide="probability of P-value <  $\alpha$ ")
47     s = (a < 0 ? "-" : "+") * string(round(a; digits=4))
48     title!("$(testname)X: $(distname(distx)), m=$m\n\
49           Y: $(distname(disty))$s, n=$n")
50     plot!(size=(400, 450))
51     plot!(; kwargs...)
52 end
53
54 function plot_pvals();
55     distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10,
56     L = 10^6, a = nothing, Δμ = nothing, kwargs...
57     @show (mean(distx), std(distx))
58     @show (mean(disty), std(disty))
59
60     if isnothing(a)
61         @show a = tieshift(distx, disty)
62         @show prob_x_le_y(distx, disty + a)
63     else
64         @show a
65         @show median(distx) - median(disty)
66     end
67     if isnothing(Δμ)
68         @show Δμ = mean(distx) - mean(disty)
69         @show mean(distx), mean(disty + Δμ)
70     else
71         @show Δμ
72         @show mean(distx), mean(disty + Δμ)
73     end
74
75     ecdf_bm = @time sim_brunner_mumzel();
76         distx = distx,
77         disty = disty + a,

```

```

78     m, n, L, kwargs...)
79     ecdf_w = @time sim_welch();
80     distx = distx,
81     disty = disty + Δμ,
82     m, n, L, kwargs...)
83     ymax = max(ecdf_bm(0.1), ecdf_w(0.1))
84     P1 = plot_ecdf(ecdf_bm, distx, disty, m, n, a;
85         testname="Brunner-Munzel test\n",
86         ylim=(-0.002, 1.02*ymax), kwargs...)
87     P2 = plot_ecdf(ecdf_w, distx, disty, m, n, Δμ;
88         testname="Welch t-test\n",
89         ylim=(-0.002, 1.02*ymax), kwargs...)
90     plot(P1, P2; size=(800, 450), topmargin=3.5Plots.mm)
91 end

```

Out[42]: `plot_pvals` (generic function with 1 method)

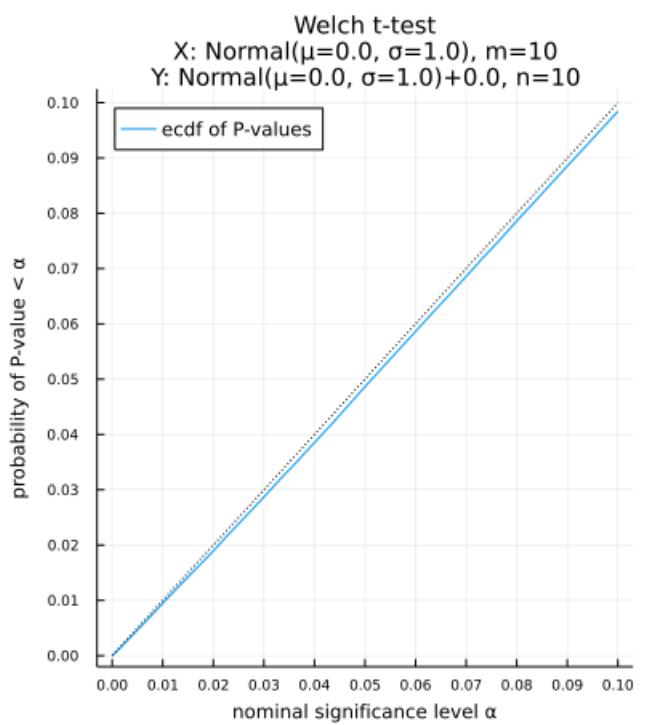
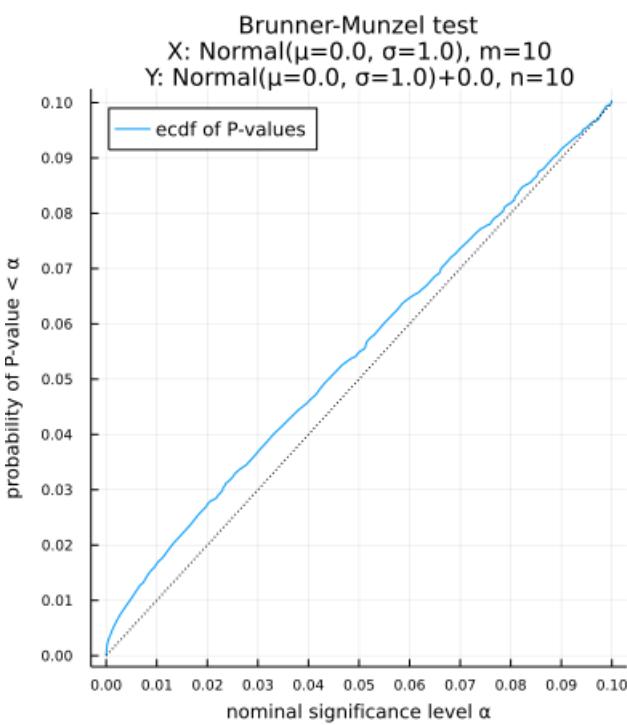
In [43]: 1 `plot_pvals(; distx = Normal(0, 1), disty = Normal(0, 1), m = 10, n = 10)`

```

(mean(distx), std(distx)) = (0.0, 1.0)
(mean(disty), std(disty)) = (0.0, 1.0)
a = tieshift(distx, disty) = 0.0
prob_x_le_y(distx, disty + a) = 0.5
Δμ = mean(distx) - mean(disty) = 0.0
(mean(distx), mean(disty + Δμ)) = (0.0, 0.0)
0.283915 seconds (216.93 k allocations: 33.653 MiB, 25.81% compilation time: 36% of which was
recompilation)
0.250943 seconds (55.78 k allocations: 25.793 MiB, 9.82% gc time, 22.42% compilation time)

```

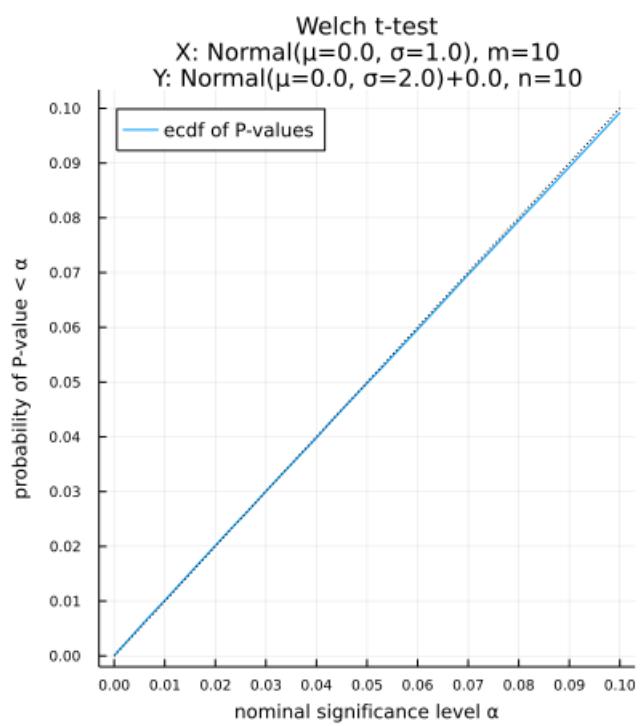
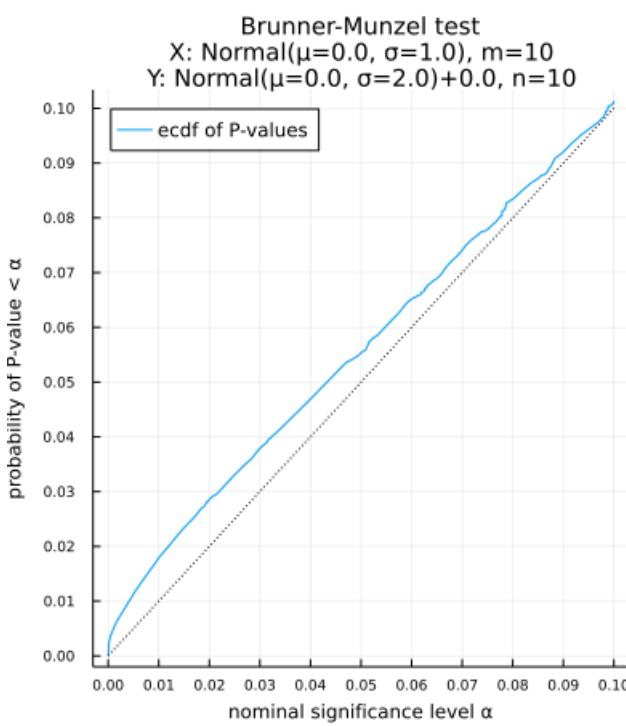
Out[43]:



```
In [44]: 1 plot_pvals(; distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10)
```

```
(mean(distx), std(distx)) = (0.0, 1.0)
(mean(disty), std(disty)) = (0.0, 2.0)
a = tieshift(distx, disty) = 7.685641860444171e-14
prob_x_le_y(distx, disty + a) = 0.5
Δμ = mean(distx) - mean(disty) = 0.0
(mean(distx), mean(disty + Δμ)) = (0.0, 0.0)
0.241331 seconds (241 allocations: 22.914 MiB)
0.163724 seconds (111 allocations: 22.900 MiB)
```

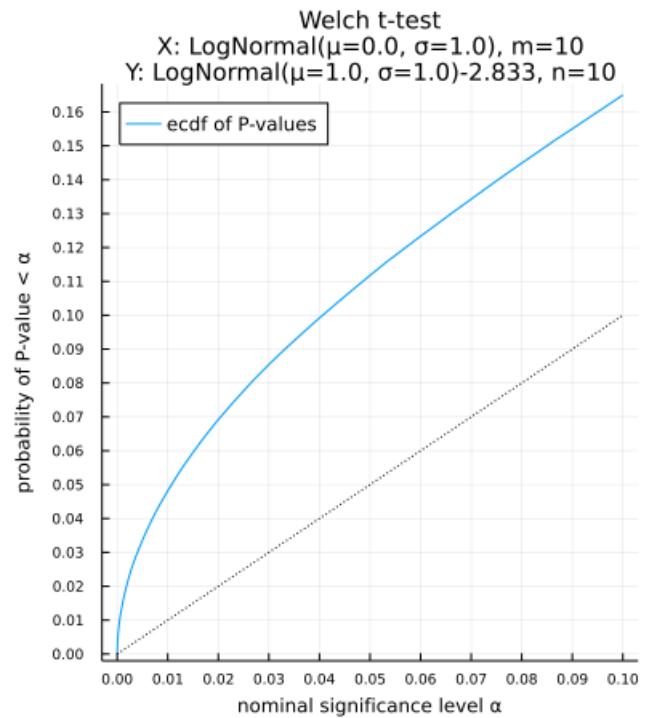
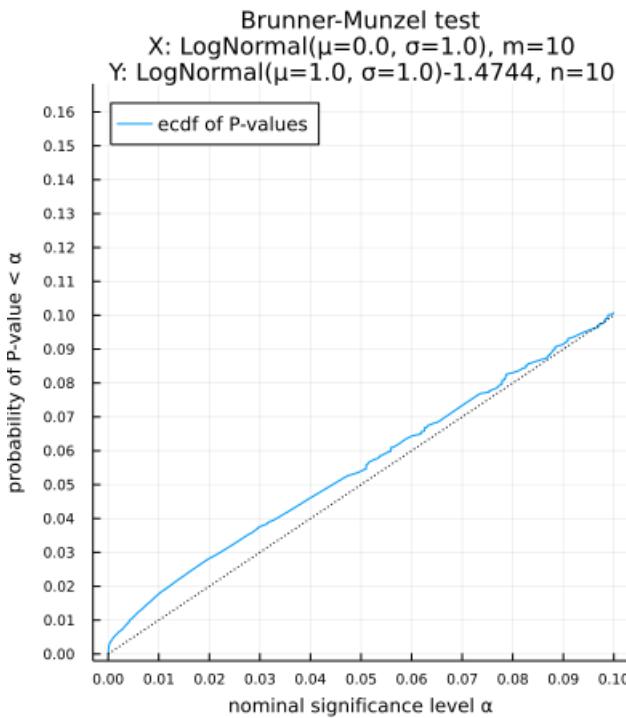
Out[44]:



```
In [45]: 1 plot_pvals(; distx = LogNormal(), disty = LogNormal(1), m = 10, n = 10)
```

```
(mean(distx), std(distx)) = (1.6487212707001282, 2.1611974158950877)
(mean(disty), std(disty)) = (4.4816890703380645, 5.874743663340262)
a = tieshift(distx, disty) = -1.4744426128871542
prob_x_le_y(distx, disty + a) = 0.5
Δμ = mean(distx) - mean(disty) = -2.8329677996379363
(mean(distx), mean(disty + Δμ)) = (1.6487212707001282, 1.6487212707001282)
0.291390 seconds (38.05 k allocations: 24.837 MiB, 12.10% compilation time)
0.268724 seconds (21.78 k allocations: 23.992 MiB, 9.71% compilation time)
```

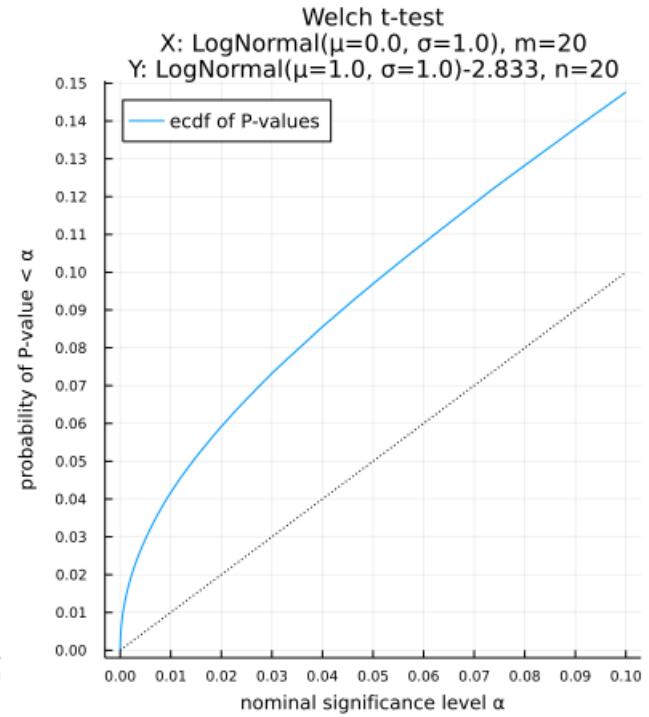
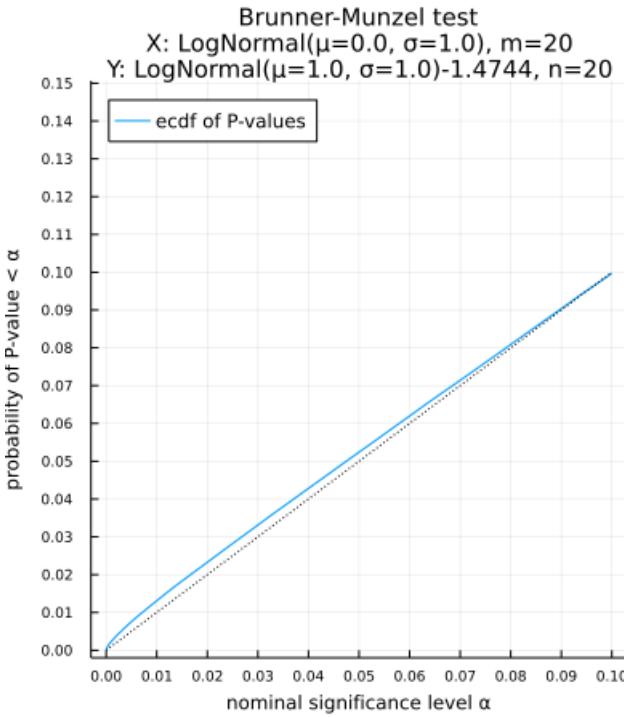
Out[45]:



```
In [46]: 1 | plot_pvals(; distx = LogNormal(), disty = LogNormal(1), m = 20, n = 20)
```

```
(mean(distx), std(distx)) = (1.6487212707001282, 2.1611974158950877)
(mean(disty), std(disty)) = (4.4816890703380645, 5.874743663340262)
a = tieshift(distx, disty) = -1.4744426128871542
prob_x_le_y(distx, disty + a) = 0.5
Δμ = mean(distx) - mean(disty) = -2.8329677996379363
(mean(distx), mean(disty + Δμ)) = (1.6487212707001282, 1.6487212707001282)
0.404015 seconds (242 allocations: 22.918 MiB)
0.249373 seconds (212 allocations: 22.911 MiB)
```

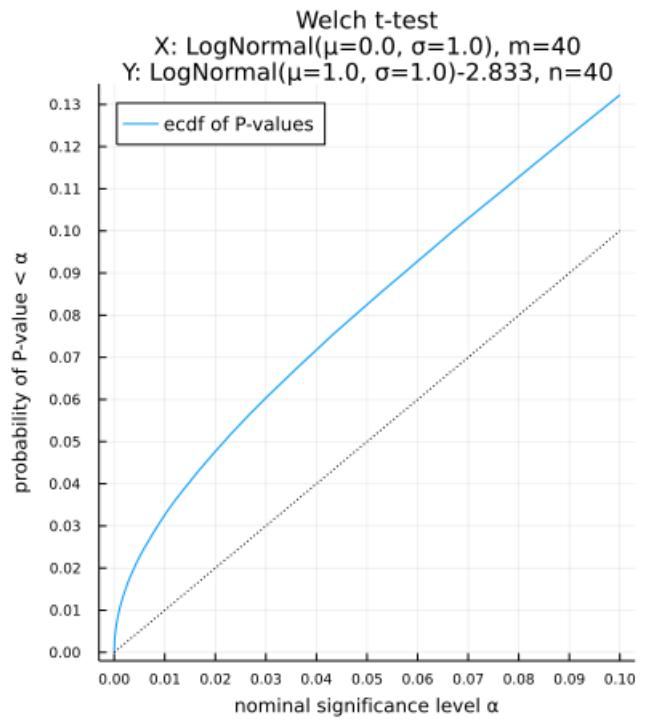
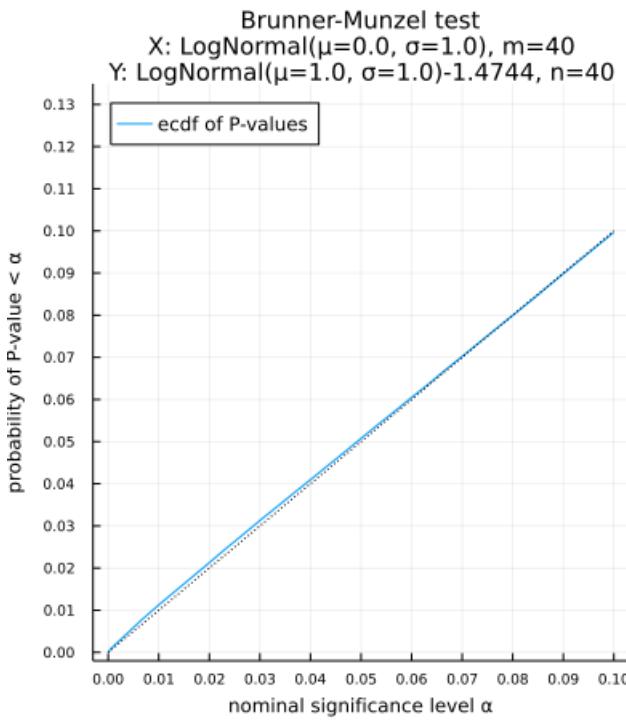
Out[46]:



```
In [47]: 1 plot_pvals(; distx = LogNormal(), disty = LogNormal(1), m = 40, n = 40)
```

```
(mean(distx), std(distx)) = (1.6487212707001282, 2.1611974158950877)
(mean(disty), std(disty)) = (4.4816890703380645, 5.874743663340262)
a = tieshift(distx, disty) = -1.4744426128871542
prob_x_le_y(distx, disty + a) = 0.5
Δμ = mean(distx) - mean(disty) = -2.8329677996379363
(mean(distx), mean(disty + Δμ)) = (1.6487212707001282, 1.6487212707001282)
0.887567 seconds (238 allocations: 22.926 MiB, 3.63% gc time)
0.413472 seconds (217 allocations: 22.915 MiB)
```

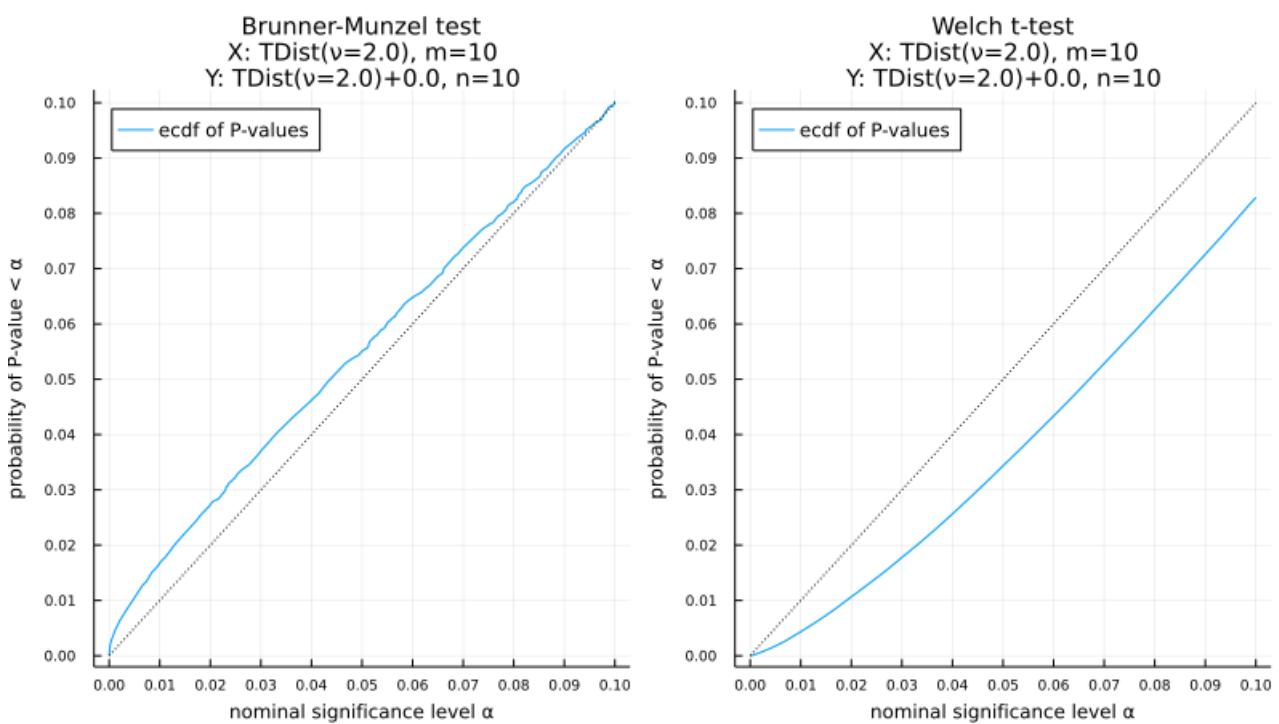
Out[47]:



```
In [48]: 1 plot_pvals(; distx = TDist(2), disty = TDist(2), m = 10, n = 10, Δμ = 0.0)
```

```
(mean(distx), std(distx)) = (0.0, Inf)
(mean(disty), std(disty)) = (0.0, Inf)
a = tieshift(distx, disty) = 0.0
prob_x_le_y(distx, disty + a) = 0.5000000000000001
Δμ = 0.0
(mean(distx), mean(disty + Δμ)) = (0.0, 0.0)
0.438389 seconds (365.61 k allocations: 41.298 MiB, 35.98% compilation time)
0.252575 seconds (21.88 k allocations: 24.050 MiB, 10.27% compilation time)
```

Out[48]:

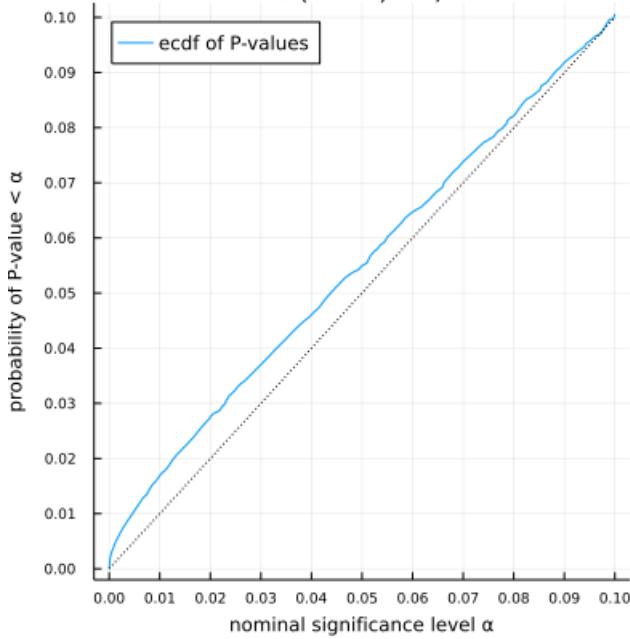


```
In [49]: 1 | plot_pvals(; distx = TDist(2), disty = TDist(1.1), m = 10, n = 10, Δμ = 0.0)
```

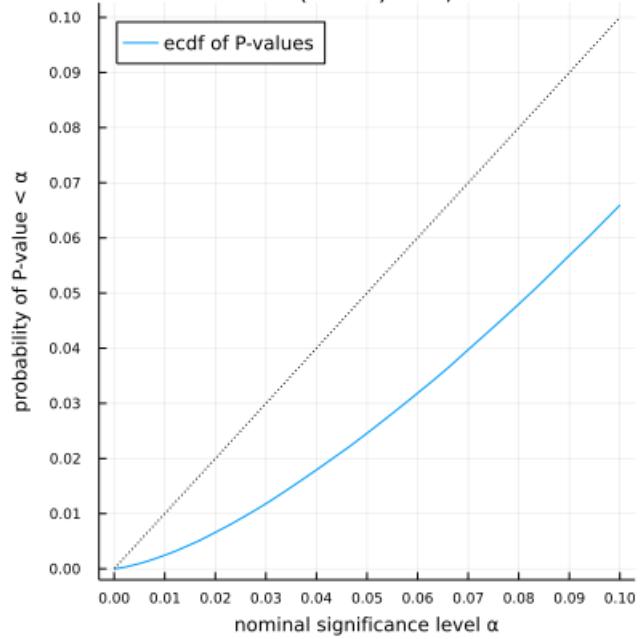
```
(mean(distx), std(distx)) = (0.0, Inf)
(mean(disty), std(disty)) = (0.0, Inf)
a = tieshift(distx, disty) = -3.4064499775914207e-9
prob_x_le_y(distx, disty + a) = 0.5
Δμ = 0.0
(mean(distx), mean(disty + Δμ)) = (0.0, 0.0)
0.410924 seconds (243 allocations: 22.914 MiB, 5.76% gc time)
0.315841 seconds (212 allocations: 22.909 MiB)
```

Out[49]:

Brunner-Munzel test
X: TDist(v=2.0), m=10
Y: TDist(v=1.1)-0.0, n=10



Welch t-test
X: TDist(v=2.0), m=10
Y: TDist(v=1.1)+0.0, n=10

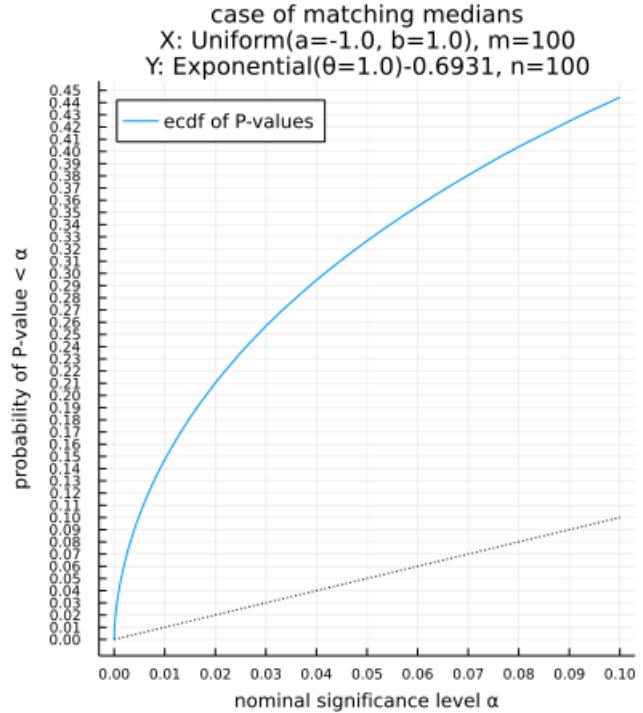
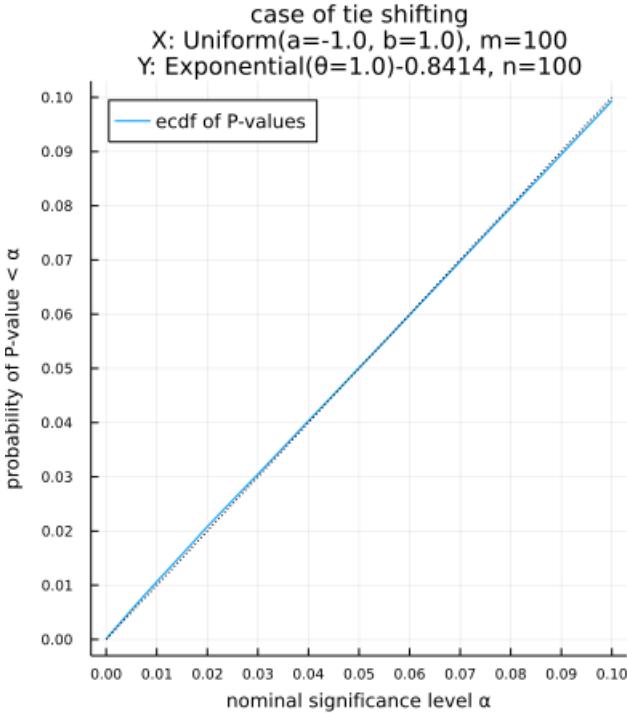


4.2 Brunner-Munzel検定は中央値に関する検定ではないことの証拠

```
In [50]: 1 distx, disty = Uniform(-1, 1), Exponential()
2 m, n, = 100, 100
3
4 @show distx, std(distx)
5 @show disty, std(disty)
6
7 @show a = tieshift(distx, disty)
8 ecdf_pval1 = @time sim_brunner_mumzel();
9     distx = distx, disty = disty + a, m, n)
10 P1 = plot_ecdf(ecdf_pval1, distx, disty, m, n, a;
11     testname="case of tie shifting\n")
12
13 @show a = median(distx) - median(disty)
14 ecdf_pval2 = @time sim_brunner_mumzel();
15     distx = distx, disty = disty + a, m, n)
16 P2 = plot_ecdf(ecdf_pval2, distx, disty, m, n, a;
17     testname="case of matching medians\n")
18
19 plot(P1, P2; size=(800, 450), topmargin=4Plots.mm)
```

```
(distx, std(distx)) = (Uniform{Float64}(a=-1.0, b=1.0), 0.5773502691896257)
(disty, std(disty)) = (Exponential{Float64}(\theta=1.0), 1.0)
a = tieshift(distx, disty) = -0.8414056600399943
3.336798 seconds (436.95 k allocations: 45.546 MiB, 7.31% compilation time)
a = median(distx) - median(disty) = -0.6931471805599453
3.105027 seconds (246 allocations: 22.947 MiB)
```

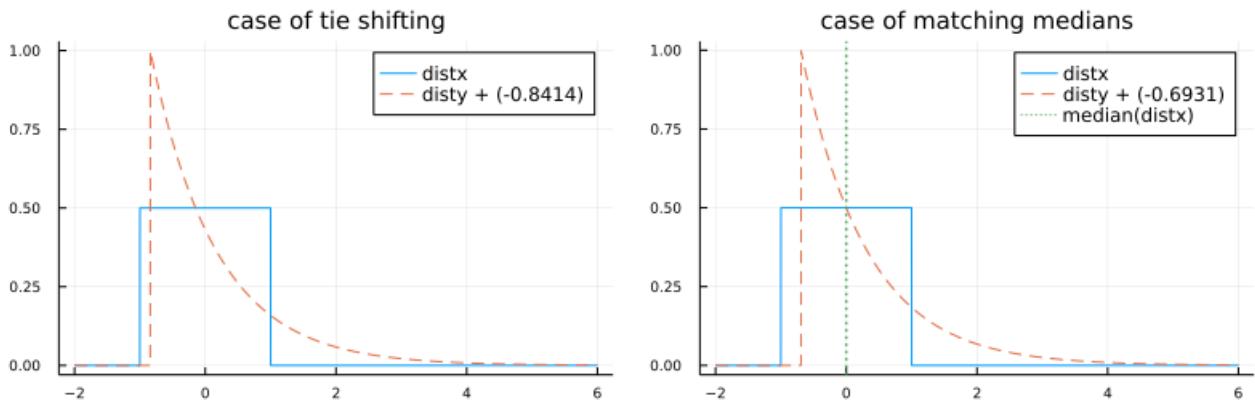
Out[50]:



```
In [51]: 1 distx, disty = Uniform(-1, 1), Exponential()
2 @show distx, std(distx)
3 @show disty, std(disty)
4
5 a = @show tieshift(distx, disty)
6 P1 = plot(distx, -2, 6; label="distx")
7 plot!(disty + a, -2, 6; label="disty + $(round(a; digits=4))", ls=:dash)
8 title!("case of tie shifting")
9
10 a = @show median(distx) - median(disty)
11 P2 = plot(distx, -2, 6; label="distx")
12 plot!(disty + a, -2, 6; label="disty + $(round(a; digits=4))", ls=:dash)
13 vline!([median(distx)]; label="median(distx)", ls=:dot, lw=1.5)
14 title!("case of matching medians")
15
16 plot(P1, P2; size=(800, 250))
```

```
(distx, std(distx)) = (Uniform{Float64}(a=-1.0, b=1.0), 0.5773502691896257)
(disty, std(disty)) = (Exponential{Float64}(\theta=1.0), 1.0)
tieshift(distx, disty) = -0.8414056600399943
median(distx) - median(disty) = -0.6931471805599453
```

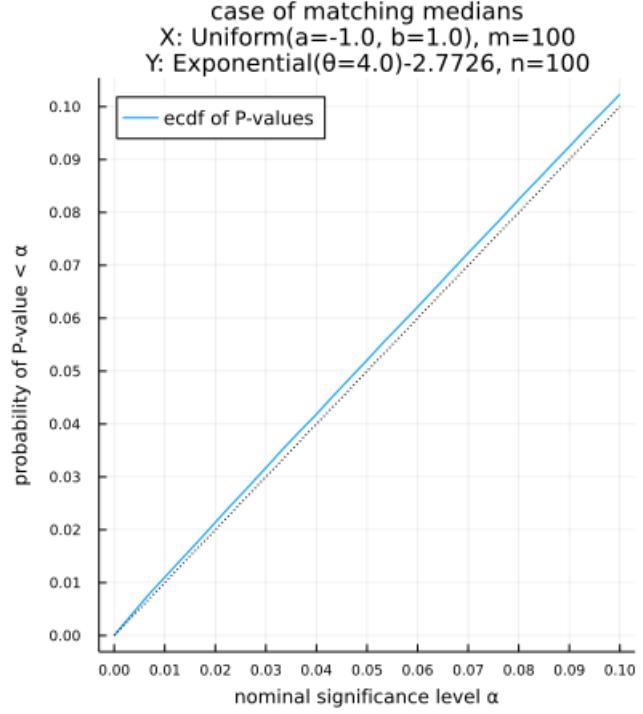
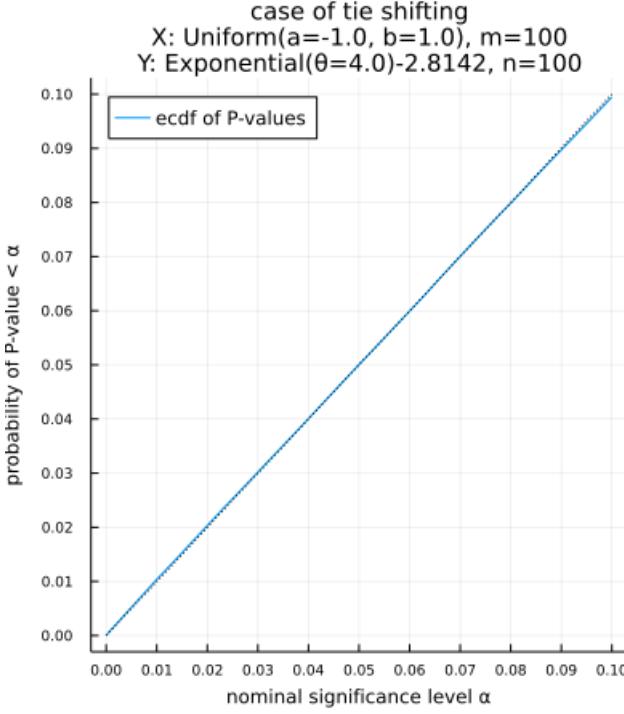
Out[51]:



```
In [52]: 1 distx, disty = Uniform(-1, 1), Exponential(4)
2 m, n, = 100, 100
3
4 @show distx, std(distx)
5 @show disty, std(disty)
6
7 @show a = tieshift(distx, disty)
8 ecdf_pval1 = @time sim_brunner_mumzel();
9     distx = distx, disty = disty + a, m, n)
10 P1 = plot_ecdf(ecdf_pval1, distx, disty, m, n, a;
11     testname="case of tie shifting\n")
12
13 @show a = median(distx) - median(disty)
14 ecdf_pval2 = @time sim_brunner_mumzel();
15     distx = distx, disty = disty + a, m, n)
16 P2 = plot_ecdf(ecdf_pval2, distx, disty, m, n, a;
17     testname="case of matching medians\n")
18
19 plot(P1, P2; size=(800, 450), topmargin=4Plots.mm)
```

```
(distx, std(distx)) = (Uniform{Float64}(a=-1.0, b=1.0), 0.5773502691896257)
(disty, std(disty)) = (Exponential{Float64}(\theta=4.0), 4.0)
a = tieshift(distx, disty) = -2.814168911097315
3.238698 seconds (251 allocations: 22.948 MiB)
a = median(distx) - median(disty) = -2.772588722239781
3.162926 seconds (238 allocations: 22.947 MiB, 0.55% gc time)
```

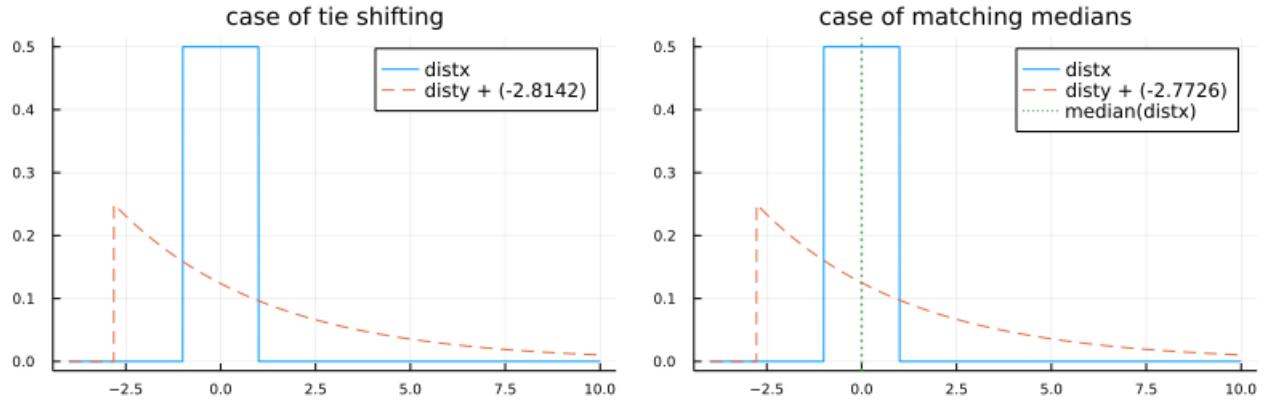
Out[52]:



```
In [53]: 1 distx, disty = Uniform(-1, 1), Exponential(4)
2 @show distx, std(distx)
3 @show disty, std(disty)
4
5 a = @show tieshift(distx, disty)
6 P1 = plot(distx, -4, 10; label="distx")
7 plot!(disty + a, -4, 10; label="disty + $(round(a; digits=4))", ls=:dash)
8 title!("case of tie shifting")
9
10 a = @show median(distx) - median(disty)
11 P2 = plot(distx, -4, 10; label="distx")
12 plot!(disty + a, -4, 10; label="disty + $(round(a; digits=4))", ls=:dash)
13 vline!([median(distx)]; label="median(distx)", ls=:dot, lw=1.5)
14 title!("case of matching medians")
15
16 plot(P1, P2; size=(800, 250))
```

(distx, std(distx)) = (Uniform{Float64}(a=-1.0, b=1.0), 0.5773502691896257)
 (disty, std(disty)) = (Exponential{Float64}($\theta=4.0$), 4.0)
 tieshift(distx, disty) = -2.814168911097315
 median(distx) - median(disty) = -2.772588722239781

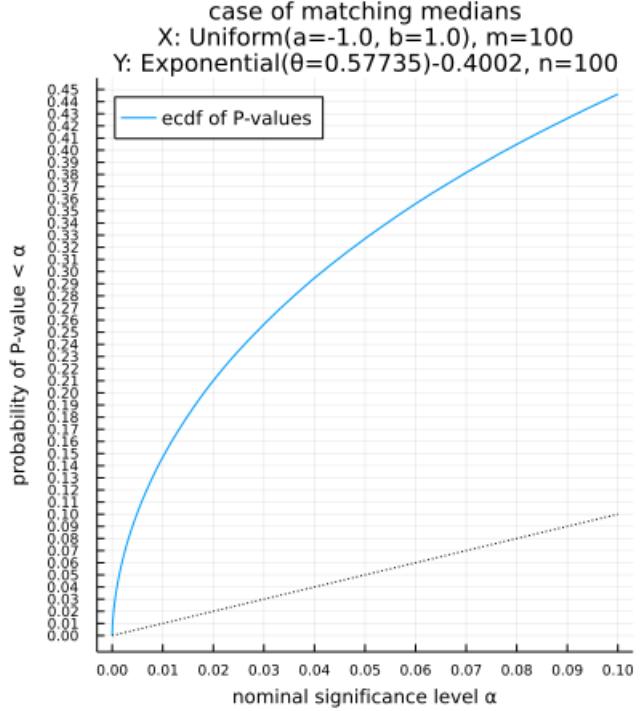
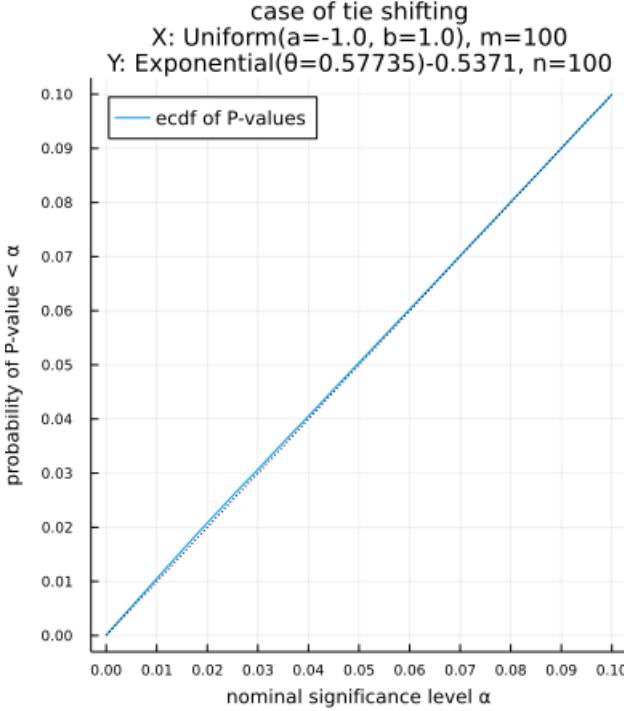
Out[53]:



```
In [54]: 1 distx, disty = Uniform(-1, 1), Exponential(0.5773502691896257)
2 m, n, = 100, 100
3
4 @show distx, std(distx)
5 @show disty, std(disty)
6
7 @show a = tieshift(distx, disty)
8 ecdf_pval1 = @time sim_brunner_mumzel();
9     distx = distx, disty = disty + a, m, n)
10 P1 = plot_ecdf(ecdf_pval1, distx, disty, m, n, a;
11     testname="case of tie shifting\n")
12
13 @show a = median(distx) - median(disty)
14 ecdf_pval2 = @time sim_brunner_mumzel();
15     distx = distx, disty = disty + a, m, n)
16 P2 = plot_ecdf(ecdf_pval2, distx, disty, m, n, a;
17     testname="case of matching medians\n")
18
19 plot(P1, P2; size=(800, 450), topmargin=4Plots.mm)
```

```
(distx, std(distx)) = (Uniform{Float64}(a=-1.0, b=1.0), 0.5773502691896257)
(disty, std(disty)) = (Exponential{Float64}(\theta=0.5773502691896257), 0.5773502691896257)
a = tieshift(distx, disty) = -0.5370568188698568
3.222829 seconds (241 allocations: 22.948 MiB)
a = median(distx) - median(disty) = -0.40018871128431455
3.177196 seconds (246 allocations: 22.948 MiB)
```

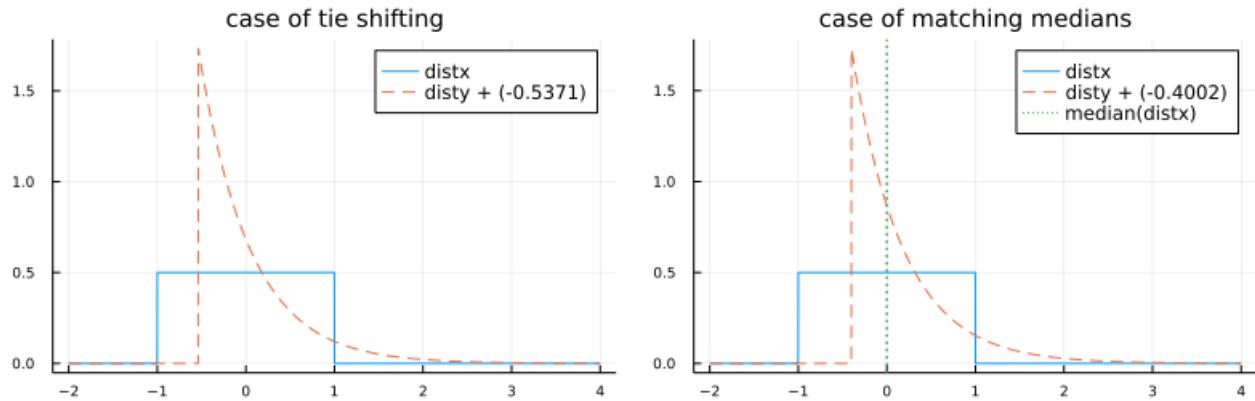
Out[54]:



```
In [55]: 1 distx, disty = Uniform(-1, 1), Exponential(0.5773502691896257)
2 @show distx, std(distx)
3 @show disty, std(disty)
4
5 a = @show tieshift(distx, disty)
6 P1 = plot(distx, -2, 4; label="distx")
7 plot!(disty + a, -2, 4; label="disty + $(round(a; digits=4))", ls=:dash)
8 title!("case of tie shifting")
9
10 a = @show median(distx) - median(disty)
11 P2 = plot(distx, -2, 4; label="distx")
12 plot!(disty + a, -2, 4; label="disty + $(round(a; digits=4))", ls=:dash)
13 vline!([median(distx)]; label="median(distx)", ls=:dot, lw=1.5)
14 title!("case of matching medians")
15
16 plot(P1, P2; size=(800, 250))
```

```
(distx, std(distx)) = (Uniform{Float64}(a=-1.0, b=1.0), 0.5773502691896257)
(disty, std(disty)) = (Exponential{Float64}(@θ=0.5773502691896257), 0.5773502691896257)
tieshift(distx, disty) = -0.5370568188698568
median(distx) - median(disty) = -0.40018871128431455
```

Out[55]:



4.3 BM検定による互角シフトの信頼区間とWelchのt検定による平均の差の信頼区間の比較

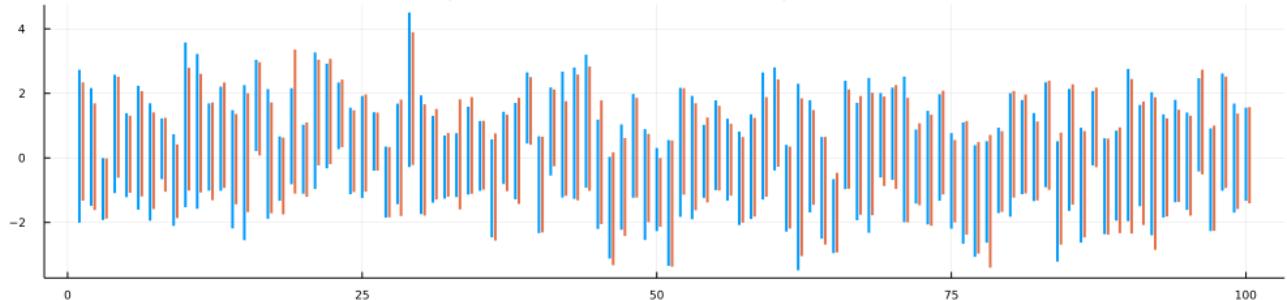
```
In [56]: 1 function plot_confints();
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10,
3     L = 100, kwargs...)
4     a = tieshift(distx, disty)
5     Δμ = mean(distx) - mean(disty)
6     BM = fill(zeros(2), 0)
7     W = fill(zeros(2), 0)
8     for _ in 1:L
9         X = rand(distx, m)
10        Y = rand(disty, n)
11        push!(BM, brunner_munzel(X, Y .+ a).confint_shift)
12        push!(W, confint_welch(X, Y .+ Δμ))
13    end
14    P = plot()
15    for i in 1:L
16        plot!(fill(i, 2), [first(BM[i]), last(BM[i])]; label="", c=1, lw=2)
17        plot!(fill(i+0.3, 2), [first(W[i]), last(W[i])]; label="", c=2, lw=2)
18    end
19    title!("X: $(distname(distx)), m=$m, Y: $(distname(disty)), n=$n")
20    plot!(size=(1000, 250))
21 end
```

Out[56]: plot_confints (generic function with 1 method)

```
In [57]: 1 plot_confints(distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10)
```

Out[57]:

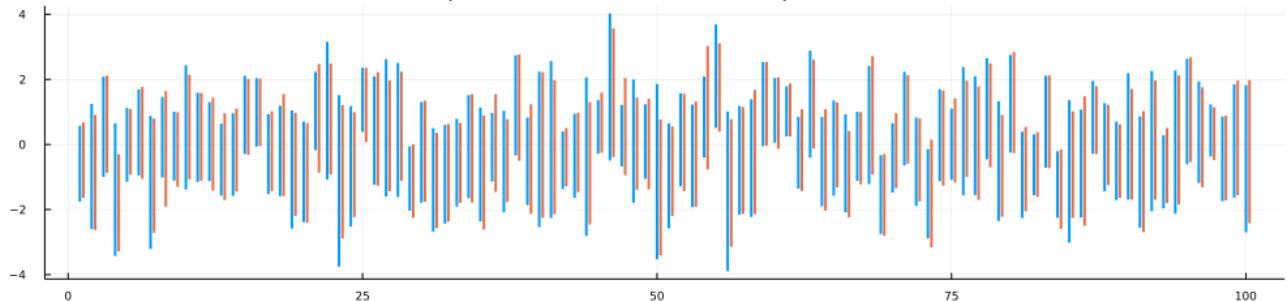
X: Normal($\mu=0.0, \sigma=1.0$), m=10, Y: Normal($\mu=0.0, \sigma=2.0$), n=10



```
In [58]: 1 plot_confints(distx = Normal(2, 1), disty = Normal(0, 2), m = 10, n = 10)
```

Out[58]:

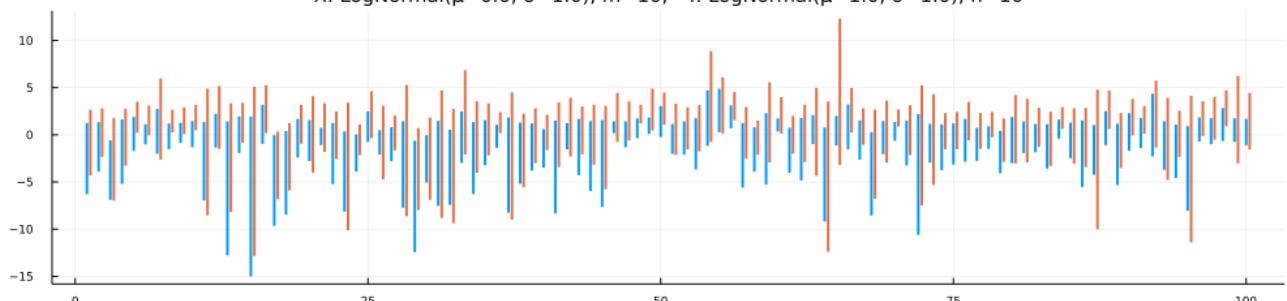
X: Normal($\mu=2.0, \sigma=1.0$), m=10, Y: Normal($\mu=0.0, \sigma=2.0$), n=10



```
In [59]: 1 plot_confints(distx = LogNormal(0), disty = LogNormal(1), m = 10, n = 10)
```

Out[59]:

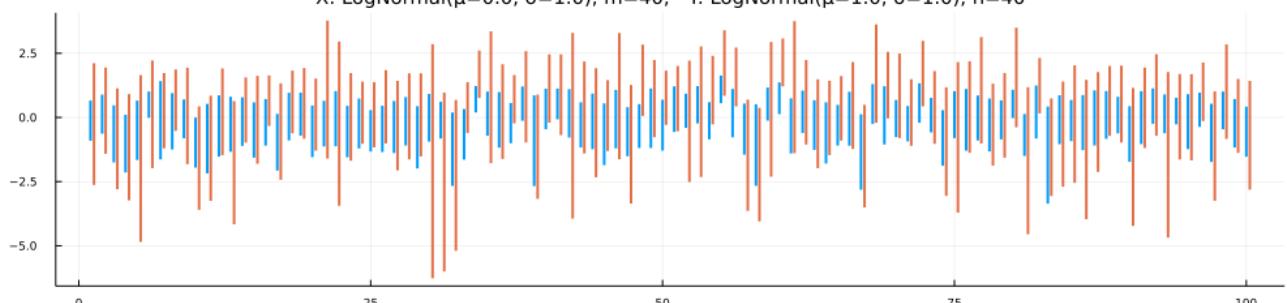
X: LogNormal($\mu=0.0, \sigma=1.0$), m=10, Y: LogNormal($\mu=1.0, \sigma=1.0$), n=10



```
In [60]: 1 plot_confints(distx = LogNormal(0), disty = LogNormal(1), m = 40, n = 40)
```

Out[60]:

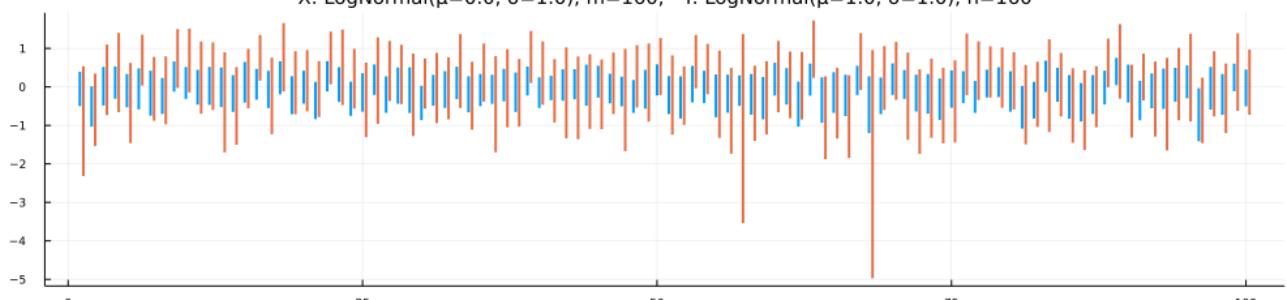
X: LogNormal($\mu=0.0, \sigma=1.0$), m=40, Y: LogNormal($\mu=1.0, \sigma=1.0$), n=40



```
In [61]: 1 plot_confints(distx = LogNormal(0), disty = LogNormal(1), m = 160, n = 160)
```

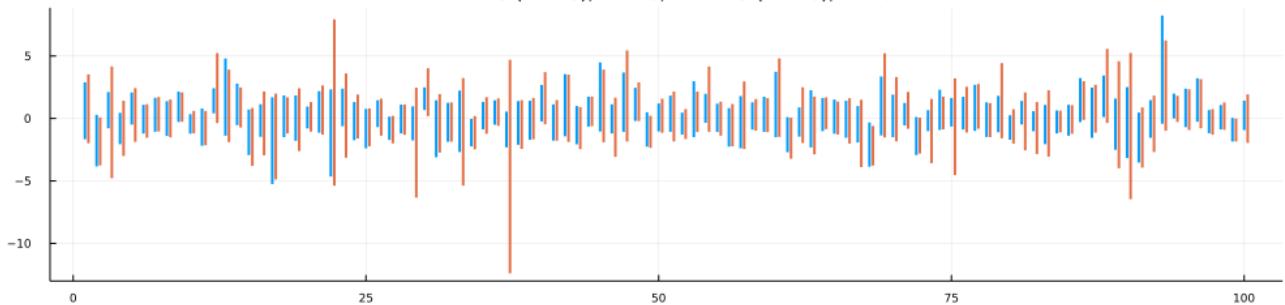
Out[61]:

X: LogNormal($\mu=0.0, \sigma=1.0$), m=160, Y: LogNormal($\mu=1.0, \sigma=1.0$), n=160



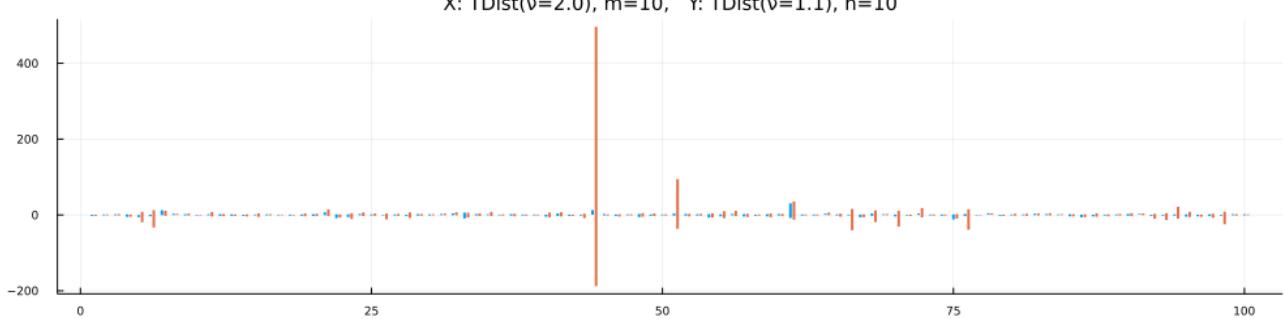
```
In [62]: 1 plot_confints(distx = TDist(2), disty = TDist(2), m = 10, n = 10)
```

Out[62]:



```
In [63]: 1 plot_confints(distx = TDist(2), disty = TDist(1.1), m = 10, n = 10)
```

Out[63]:

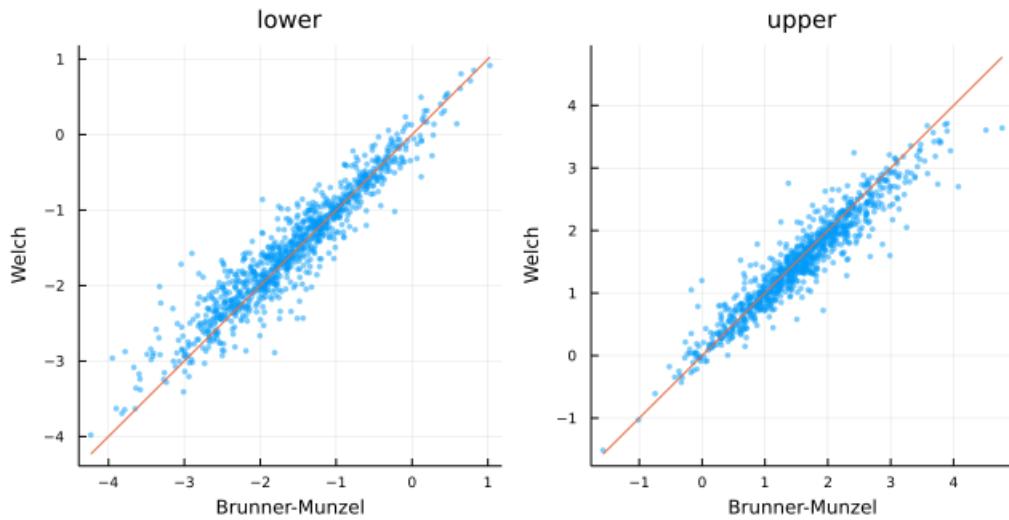


```
In [64]: 1 function plot_limits();
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10,
3     L = 1000, kwargs...
4
5     @show distx, m
6     @show disty, n
7
8     a = tieshift(distx, disty)
9     Δμ = mean(distx) - mean(disty)
10
11    BM = fill(zeros(2), 0)
12    W = fill(zeros(2), 0)
13    for _ in 1:L
14        X = rand(distx, m)
15        Y = rand(disty, n)
16        push!(BM, brunner_munzel(X, Y .+ a).confint_shift)
17        push!(W, confint_welch(X, Y .+ Δμ))
18    end
19
20    lower = [(first(BM[i]), first(W[i])) for i in 1:L]
21    upper = [(last(BM[i]), last(W[i])) for i in 1:L]
22
23    P1 = scatter(lower; label="", msc=:auto, ms=2, ma=0.5)
24    plot!(identity; label="")
25    plot!(xguide="Brunner-Munzel", yguide="Welch")
26    title!("lower")
27
28    P2 = scatter(upper; label="", msc=:auto, ms=2, ma=0.5)
29    plot!(identity; label="")
30    plot!(xguide="Brunner-Munzel", yguide="Welch")
31    title!("upper")
32
33    plot(P1, P2; size=(640, 320))
34 end
```

Out[64]: plot_limits (generic function with 1 method)

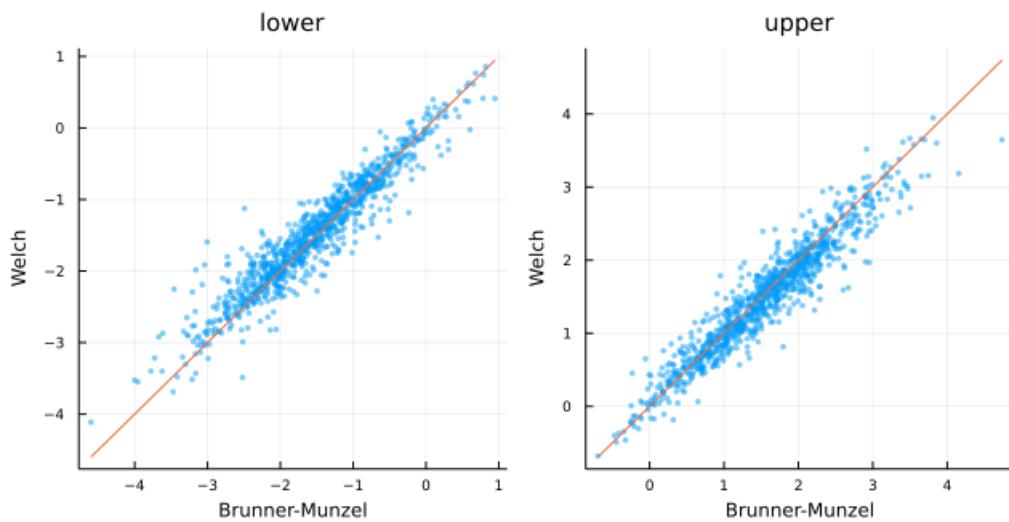
```
In [65]: 1 plot_limits(distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10)
          (distx, m) = (Normal{Float64}(\mu=0.0, σ=1.0), 10)
          (disty, n) = (Normal{Float64}(\mu=0.0, σ=2.0), 10)
```

Out[65]:



```
In [66]: 1 plot_limits(distx = Normal(2, 1), disty = Normal(0, 2), m = 10, n = 10)
          (distx, m) = (Normal{Float64}(\mu=2.0, σ=1.0), 10)
          (disty, n) = (Normal{Float64}(\mu=0.0, σ=2.0), 10)
```

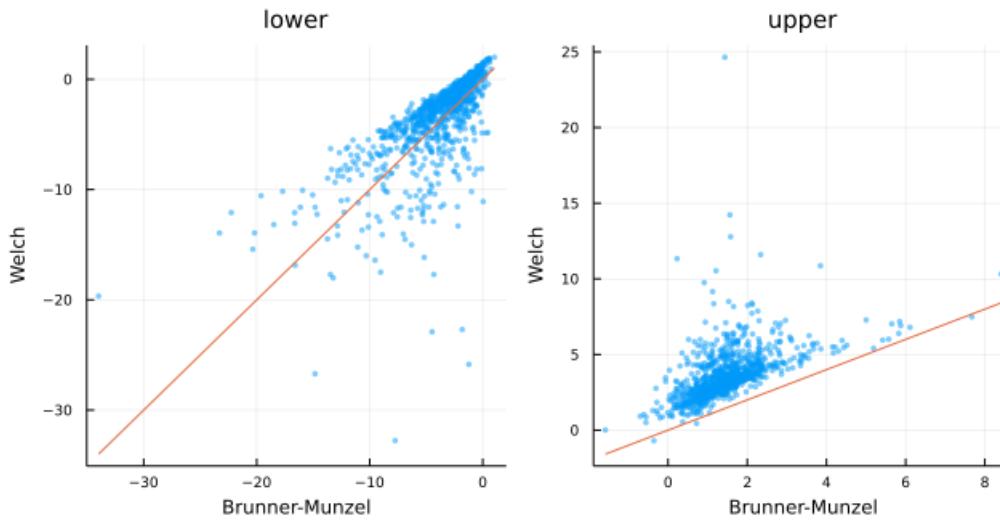
Out[66]:



```
In [67]: 1 plot_limits(distx = LogNormal(), disty = LogNormal(1), m = 10, n = 10)
```

```
(distx, m) = (LogNormal{Float64}( $\mu=0.0$ ,  $\sigma=1.0$ ), 10)
(disty, n) = (LogNormal{Float64}( $\mu=1.0$ ,  $\sigma=1.0$ ), 10)
```

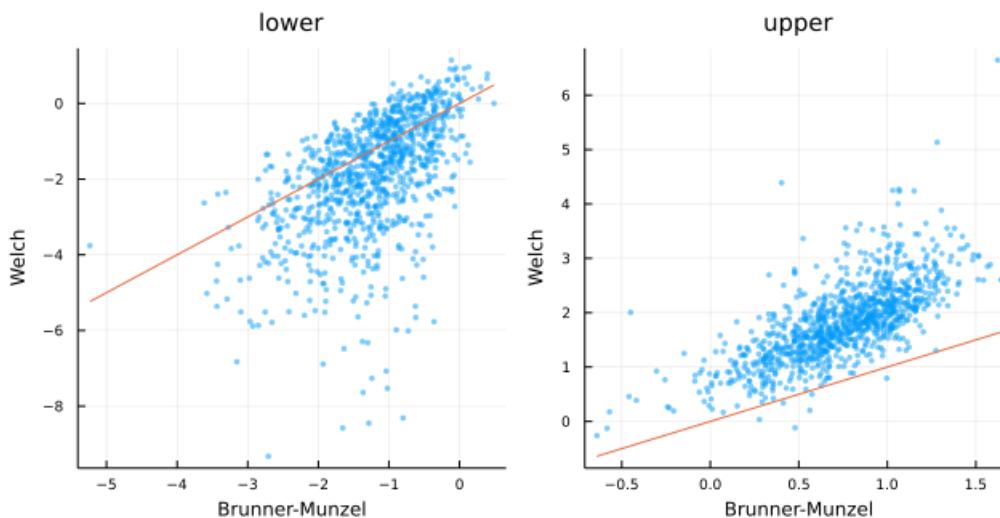
Out[67]:



```
In [68]: 1 plot_limits(distx = LogNormal(), disty = LogNormal(1), m = 40, n = 40)
```

```
(distx, m) = (LogNormal{Float64}( $\mu=0.0$ ,  $\sigma=1.0$ ), 40)
(disty, n) = (LogNormal{Float64}( $\mu=1.0$ ,  $\sigma=1.0$ ), 40)
```

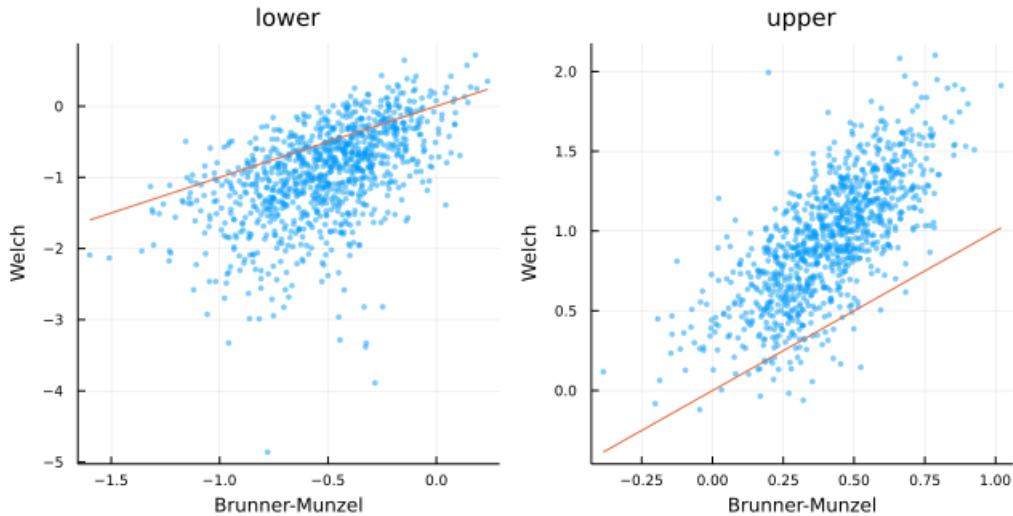
Out[68]:



```
In [69]: 1 @time plot_limits(distx = LogNormal(), disty = LogNormal(1), m = 160, n = 160)
```

```
(distx, m) = (LogNormal{Float64}( $\mu=0.0$ ,  $\sigma=1.0$ ), 160)
(disty, n) = (LogNormal{Float64}( $\mu=1.0$ ,  $\sigma=1.0$ ), 160)
6.448462 seconds (45.97 k allocations: 38.011 MiB, 0.31% gc time)
```

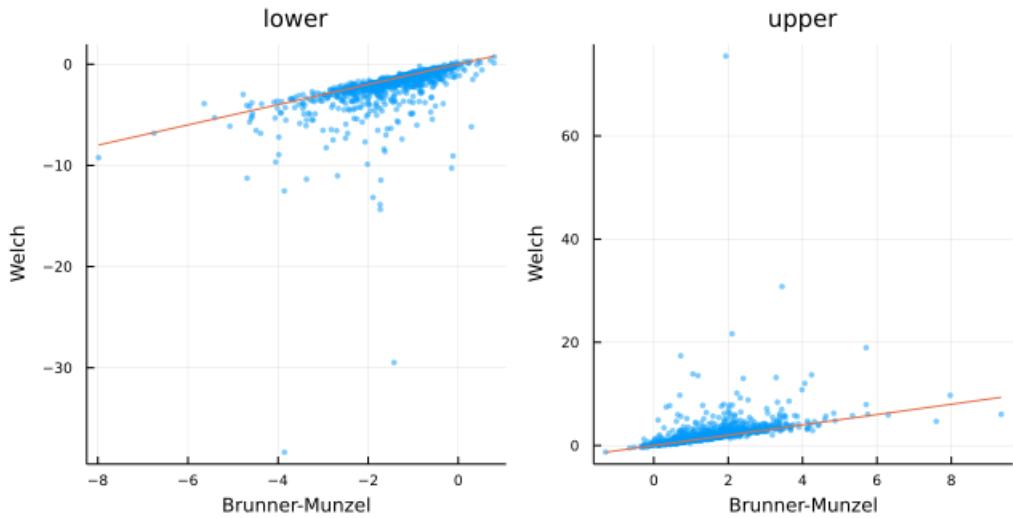
Out[69]:



```
In [70]: 1 plot_limits(distx = TDist(2), disty = TDist(2), m = 10, n = 10)
```

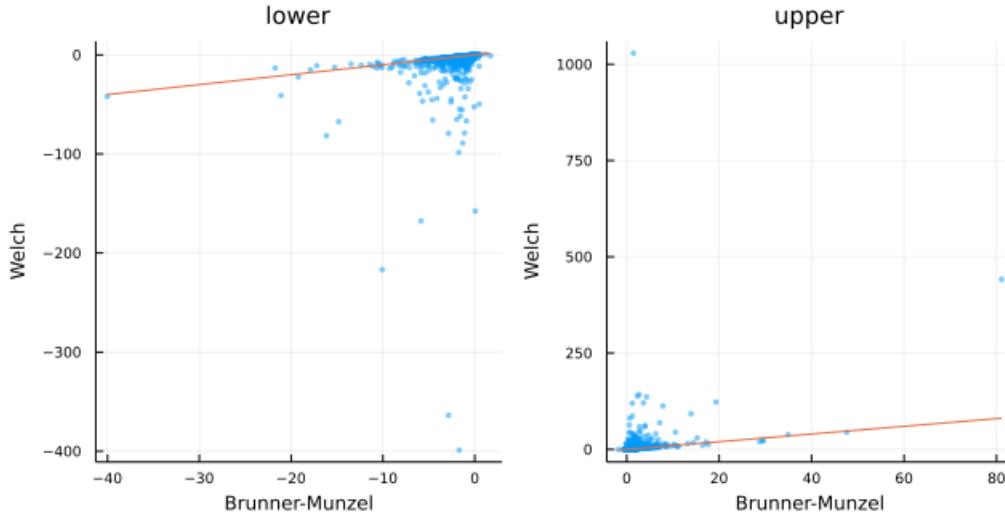
```
(distx, m) = (TDist{Float64}( $v=2.0$ ), 10)
(disty, n) = (TDist{Float64}( $v=2.0$ ), 10)
```

Out[70]:



```
In [71]: 1 plot_limits(distx = TDist(2), disty = TDist(1.1), m = 10, n = 10)
          (distx, m) = (TDist{Float64}(v=2.0), 10)
          (disty, n) = (TDist{Float64}(v=1.1), 10)
```

Out[71]:



5 小サンプルでのpermutation版の検定とBM検定とWelchのt検定の比較

```
In [72]: 1 function sim_brunner_mumzel_perm();
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 5, n = 5,
3     L = 10^2)
4     pval_bm_perm = Vector{Float64}(undef, L)
5     tmpX = [Vector{Float64}(undef, m) for _ in 1:nthreads()]
6     tmpY = [Vector{Float64}(undef, n) for _ in 1:nthreads()]
7     tmpXandY = [Vector{Float64}(undef, m+n) for _ in 1:nthreads()]
8     tmpTval = [Vector{Float64}(undef, binomial(m+n, m)) for _ in 1:nthreads()]
9     tmpHx = [Vector{Float64}(undef, m) for _ in 1:nthreads()]
10    tmpHy = [Vector{Float64}(undef, n) for _ in 1:nthreads()]
11    tmpccomb = [Vector{Int}(undef, n) for _ in 1:nthreads()]
12    @threads for i in 1:L
13        tid = threadid()
14        X = rand!(distx, tmpX[tid])
15        Y = rand!(disty, tmpY[tid])
16        Tval = permutation_tvalues_brunner_munzel(X, Y,
17            tmpXandY[tid], tmpTval[tid], tmpHx[tid], tmpHy[tid], tmpccomb[tid])
18        tval = statistics_brunner_munzel(X, Y, tmpHx[tid], tmpHy[tid]).tvalue
19        pval_bm_perm[i] = pvalue_brunner_munzel_perm(X, Y, Tval, tval)
20    end
21    ecdf(pval_bm_perm)
22 end
```

Out[72]: sim_brunner_mumzel_perm (generic function with 1 method)

```
In [73]: 1 @time ecdf_bm_perm = sim_brunner_mumzel_perm(  
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 7, n = 7, L = 10^4)
```

3.178117 seconds (35.60 k allocations: 2.909 MiB, 1.21% compilation time)

```
In [74]: 1 function plot_pvals_with_perm();
2     distx = Normal(0, 1),
3     disty = Normal(0, 2),
4     m = 7,
5     n = 7,
6     L = 10^4,
7     kwargs...
8 )
9 a = tieshift(distx, disty)
10 @time ecdf_bm_perm = sim_brunner_mumzel_perm(; distx, disty = disty + a, m, n, L)
11 @time ecdf_bm = sim_brunner_mumzel(; distx, disty = disty + a, m, n, L)
12 Δμ = mean(distx) - mean(disty)
13 @time ecdf_w = sim_welch(; distx, disty = disty + Δμ, m, n, L)
14 @show a Δμ
15
16 plot(legend=:topleft)
17 plot!(α → ecdf_bm_perm(α), 0, 0.1; label="BM permutation")
18 plot!(α → ecdf_bm(α), 0, 0.1; label="Brunner-Munzel", ls=:dash)
19 plot!(α → ecdf_w(α), 0, 0.1; label="Welch", ls=:dashdot)
20 plot!(identity; label="", c=:black, ls=:dot)
21 plot!(xtick=0:0.01:0.1, ytick=0:0.01:1)
22 plot!(xguide="nominal significance level α",
23       yguide="probability of P-value < α")
24 a_ = string(round(a; digits=4))
25 Δμ_ = string(round(Δμ; digits=4))
26 title!("X: $(distname(distx)), m=$m\n"
27        "Y: $(distname(disty))+(a, Δμ), n=$n\n"
28        "a=$a_, Δμ=$Δμ_")
29 plot!(size=(400, 450), titlefontsize=9)
30 plot!(; kwargs...)
31 end
```

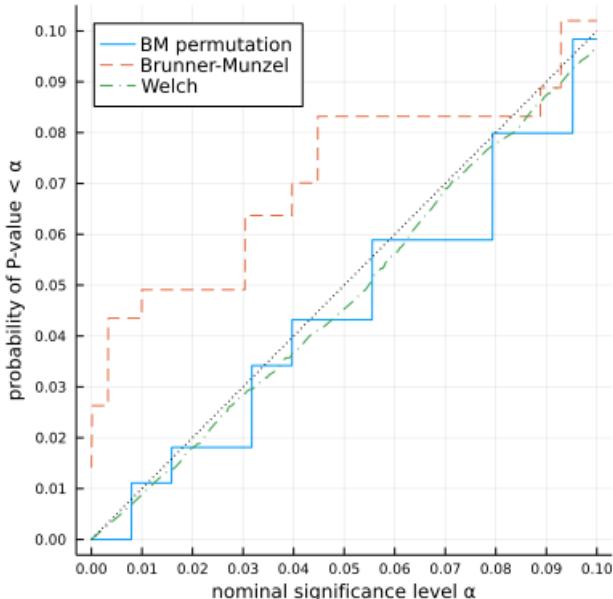
Out[74]: `plot_pvals_with_perm` (generic function with 1 method)

```
In [75]: 1 plot_pvals_with_perm(
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 5, n = 5, L = 10^4)

0.193272 seconds (10.18 k allocations: 1.186 MiB)
0.001737 seconds (144 allocations: 248.969 KiB)
0.001214 seconds (114 allocations: 245.734 KiB)
a = 7.685641860444171e-14
Δμ = 0.0
```

Out[75]:

```
X: Normal(μ=0.0, σ=1.0), m=5
Y: Normal(μ=0.0, σ=2.0)+(a, Δμ), n=5
a=0.0, Δμ=0.0
```



```
In [76]: 1 plot_pvals_with_perm()
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 7, n = 7, L = 10^4)
```

3.287972 seconds (10.19 k allocations: 1.630 MiB)

0.002010 seconds (148 allocations: 249.578 KiB)

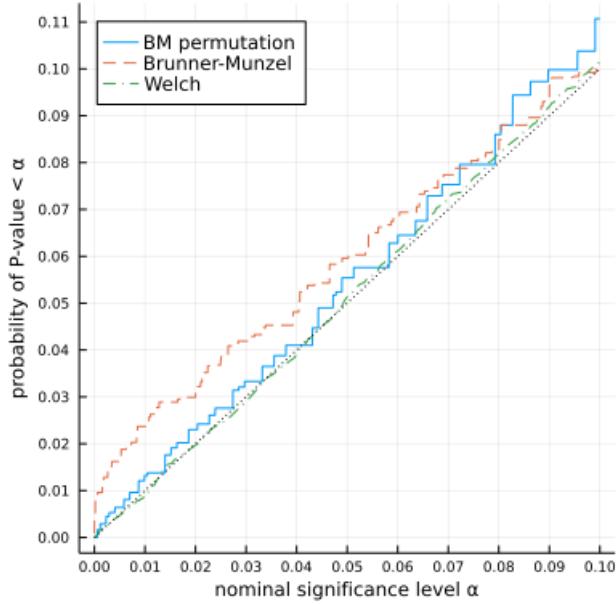
0.001511 seconds (117 allocations: 246.203 KiB)

a = 7.685641860444171e-14

$\Delta\mu = 0.0$

Out[76]:

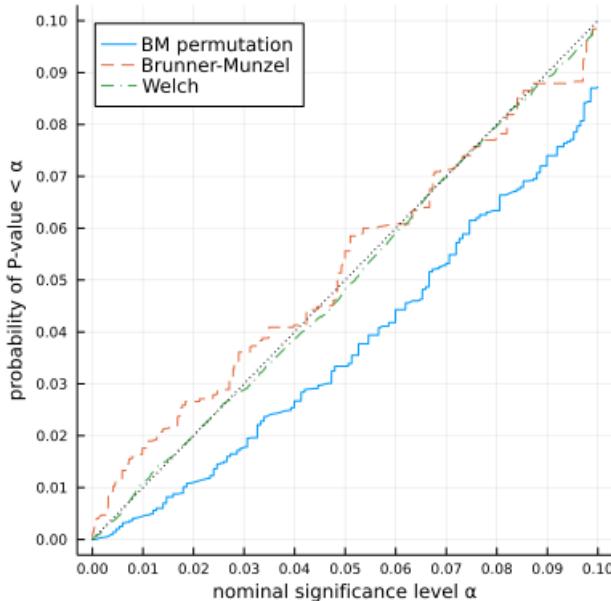
X: Normal($\mu=0.0, \sigma=1.0$), m=7
Y: Normal($\mu=0.0, \sigma=2.0$)+(a, $\Delta\mu$), n=7
a=0.0, $\Delta\mu=0.0$



```
In [77]: 1 plot_pvals_with_perm()
2         distx = Normal(0, 1), disty = Normal(0, 2), m = 5, n = 10, L = 10^4)
```

3.163012 seconds (10.20 k allocations: 1.440 MiB)
0.001998 seconds (135 allocations: 249.422 KiB)
0.001645 seconds (113 allocations: 246.266 KiB)
a = 7.685641860444171e-14
 $\Delta\mu = 0.0$

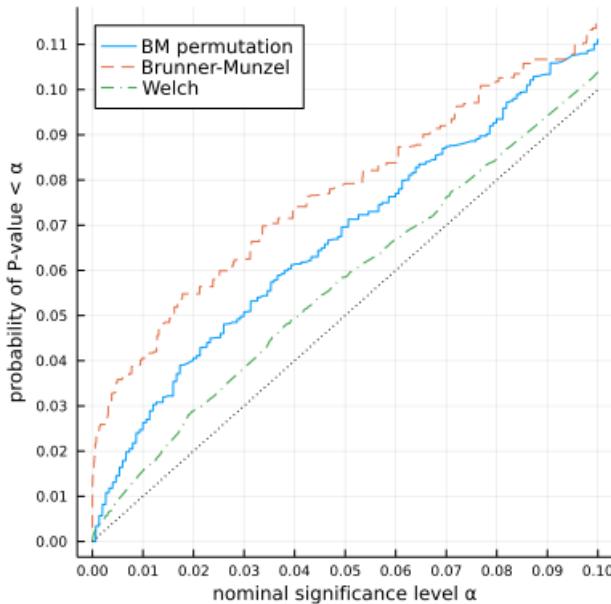
Out[77]: X: Normal($\mu=0.0, \sigma=1.0$), m=5
Y: Normal($\mu=0.0, \sigma=2.0+(a, \Delta\mu)$, n=10
a=0.0, $\Delta\mu=0.0$



```
In [78]: 1 plot_pvals_with_perm()
2         distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 5, L = 10^4)
```

3.293527 seconds (10.20 k allocations: 1.897 MiB)
0.001594 seconds (138 allocations: 249.516 KiB)
0.021377 seconds (116 allocations: 246.312 KiB)
a = 7.685641860444171e-14
 $\Delta\mu = 0.0$

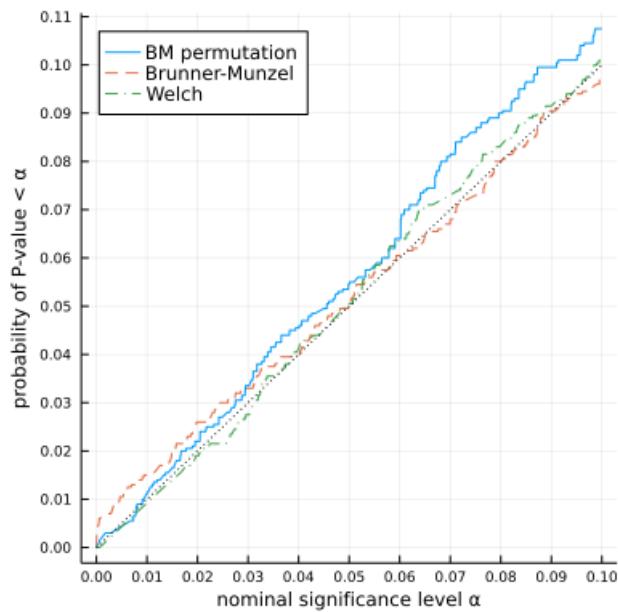
Out[78]: X: Normal($\mu=0.0, \sigma=1.0$), m=10
Y: Normal($\mu=0.0, \sigma=2.0+(a, \Delta\mu)$, n=5
a=0.0, $\Delta\mu=0.0$



```
In [79]: 1 plot_pvals_with_perm(  
2     distx = Normal(0, 1), disty = Normal(0, 2), m = 10, n = 10, L = 2000)
```

```
46.360802 seconds (2.22 k allocations: 17.258 MiB)  
0.000547 seconds (143 allocations: 63.750 KiB)  
0.019471 seconds (114 allocations: 59.906 KiB)  
a = 7.685641860444171e-14  
 $\Delta\mu = 0.0$ 
```

```
Out[79]: X: Normal( $\mu=0.0, \sigma=1.0$ ), m=10  
Y: Normal( $\mu=0.0, \sigma=2.0$ )+(a,  $\Delta\mu$ ), n=10  
a=0.0,  $\Delta\mu=0.0$ 
```



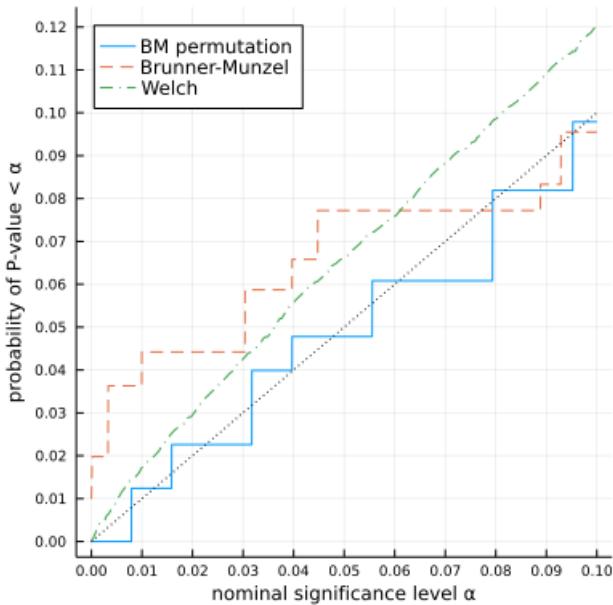
In [80]:

```
1 plot_pvals_with_perm()
2     distx = Exponential(1), disty = Exponential(2),
3     m = 5, n = 5, L = 10^4)
```

```
0.247603 seconds (40.78 k allocations: 2.758 MiB, 17.10% compilation time)
0.052518 seconds (20.11 k allocations: 1.258 MiB, 96.83% compilation time)
0.063320 seconds (18.60 k allocations: 1.199 MiB, 96.50% compilation time)
a = -0.5753641445892759
Δμ = -1.0
```

Out[80]:

X: Exponential($\theta=1.0$), m=5
Y: Exponential($\theta=2.0$)+(a, $\Delta\mu$), n=5
a=-0.5754, $\Delta\mu=-1.0$



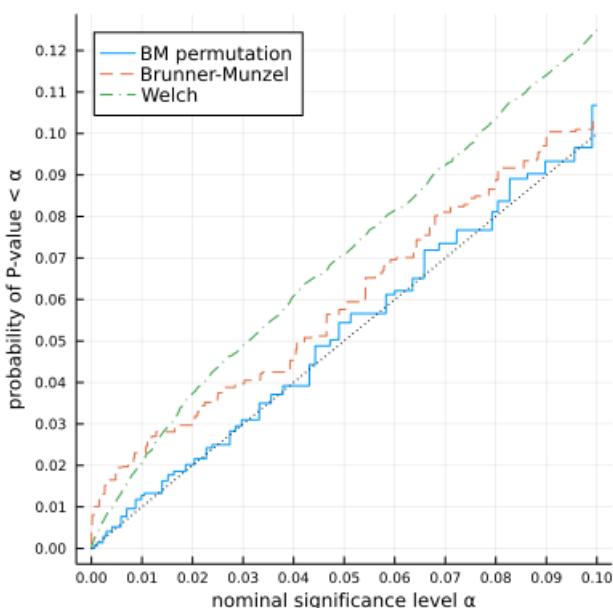
In [81]:

```
1 plot_pvals_with_perm()
2     distx = Exponential(1), disty = Exponential(2),
3     m = 7, n = 7, L = 10^4)
```

```
3.418676 seconds (10.19 k allocations: 1.630 MiB)
0.017156 seconds (145 allocations: 249.453 KiB)
0.015168 seconds (117 allocations: 246.328 KiB)
a = -0.5753641445892759
Δμ = -1.0
```

Out[81]:

X: Exponential($\theta=1.0$), m=7
Y: Exponential($\theta=2.0$)+(a, $\Delta\mu$), n=7
a=-0.5754, $\Delta\mu=-1.0$



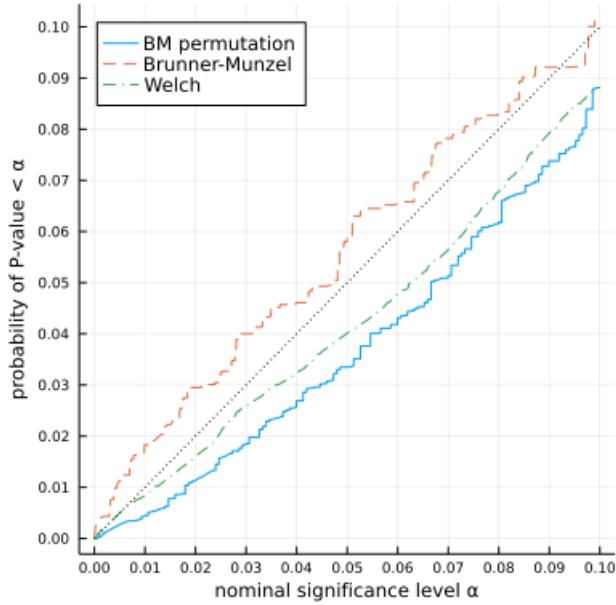
In [82]:

```
1 plot_pvals_with_perm(  
2     distx = Exponential(1), disty = Exponential(2),  
3     m = 5, n = 10, L = 10^4)
```

```
3.031049 seconds (10.19 k allocations: 1.439 MiB)  
0.001551 seconds (135 allocations: 249.422 KiB)  
0.001160 seconds (112 allocations: 246.234 KiB)  
a = -0.5753641445892759  
 $\Delta\mu = -1.0$ 
```

Out[82]:

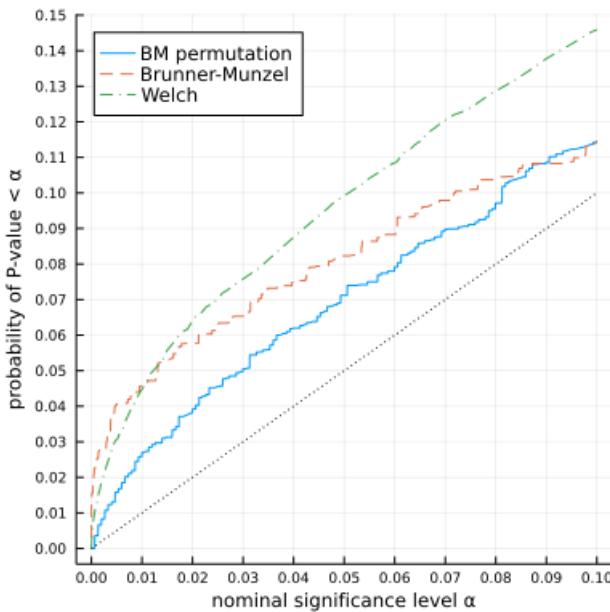
X: Exponential($\theta=1.0$), m=5
Y: Exponential($\theta=2.0$)+(a, $\Delta\mu$), n=10
a=-0.5754, $\Delta\mu=-1.0$



```
In [83]: 1 plot_pvals_with_perm()
2     distx = Exponential(1), disty = Exponential(2),
3     m = 10, n = 5, L = 10^4)
```

3.195528 seconds (10.19 k allocations: 1.897 MiB)
0.002077 seconds (145 allocations: 250.125 KiB)
0.001493 seconds (117 allocations: 246.344 KiB)
a = -0.5753641445892759
 $\Delta\mu = -1.0$

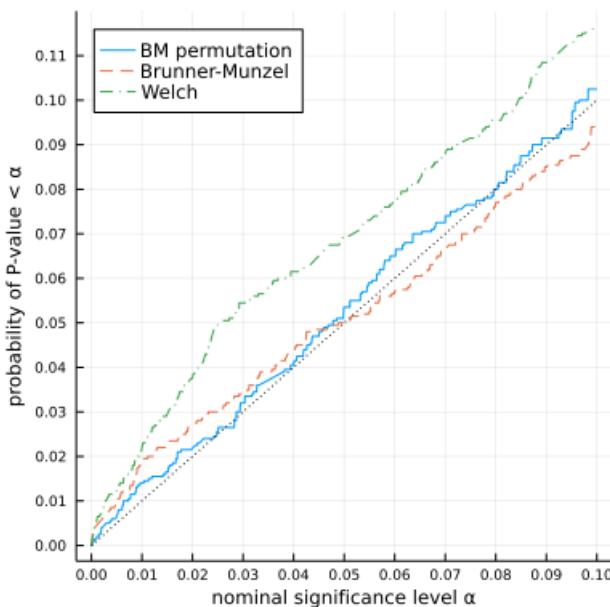
Out[83]: X: Exponential($\theta=1.0$), m=10
Y: Exponential($\theta=2.0$)+(a, $\Delta\mu$), n=5
a=-0.5754, $\Delta\mu=-1.0$



```
In [84]: 1 plot_pvals_with_perm()
2     distx = Exponential(1), disty = Exponential(2),
3     m = 10, n = 10, L = 2000)
```

47.990279 seconds (2.23 k allocations: 17.258 MiB)
0.000467 seconds (141 allocations: 63.688 KiB)
0.014886 seconds (113 allocations: 59.609 KiB)
a = -0.5753641445892759
 $\Delta\mu = -1.0$

Out[84]: X: Exponential($\theta=1.0$), m=10
Y: Exponential($\theta=2.0$)+(a, $\Delta\mu$), n=10
a=-0.5754, $\Delta\mu=-1.0$



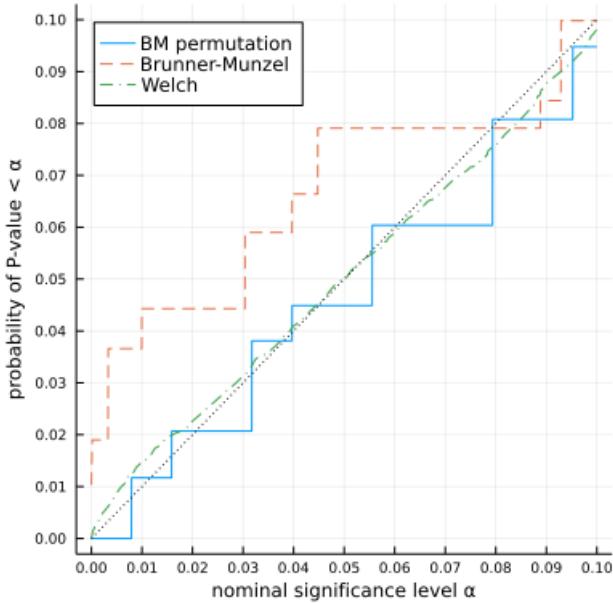
In [85]:

```
1 plot_pvals_with_perm()
2     distx = Uniform(-1, 1), disty = Exponential(0.5773502691896257),
3     m = 5, n = 5, L = 10^4)
```

```
0.255353 seconds (32.12 k allocations: 2.314 MiB, 13.21% compilation time)
0.001413 seconds (138 allocations: 248.391 KiB)
0.023070 seconds (18.50 k allocations: 1.188 MiB, 95.15% compilation time)
a = -0.5370568188698568
Δμ = -0.5773502691896257
```

Out[85]:

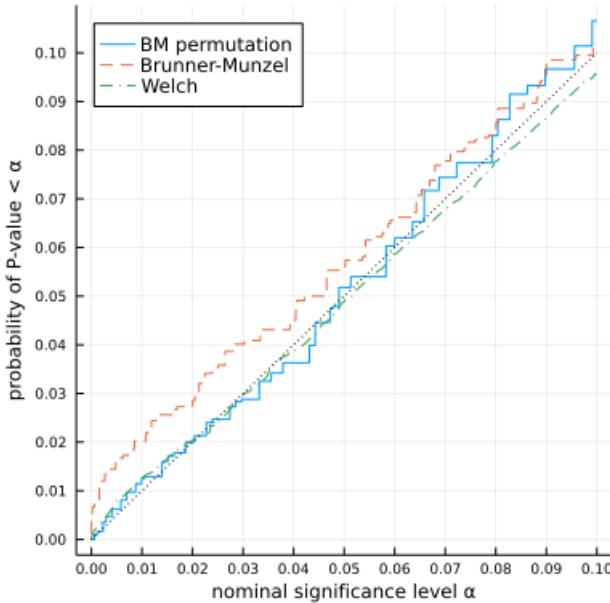
X: Uniform(a=-1.0, b=1.0), m=5
Y: Exponential($\theta=0.57735$) $+(a, \Delta\mu)$, n=5
 $a=-0.5371, \Delta\mu=-0.5774$



```
In [86]: 1 plot_pvals_with_perm()
2     distx = Uniform(-1, 1), disty = Exponential(0.5773502691896257),
3     m = 7, n = 7, L = 10^4)
```

3.419962 seconds (10.19 k allocations: 1.630 MiB)
0.025878 seconds (143 allocations: 249.422 KiB)
0.005994 seconds (118 allocations: 246.188 KiB)
 $a = -0.5370568188698568$
 $\Delta\mu = -0.5773502691896257$

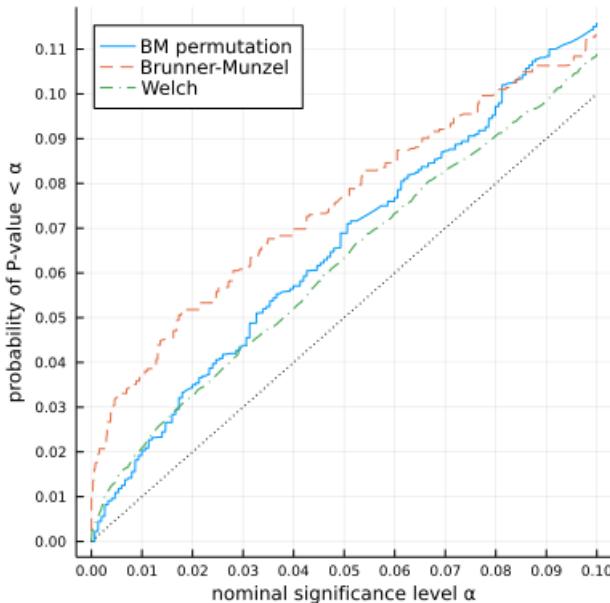
Out[86]: X: Uniform(a=-1.0, b=1.0), m=7
Y: Exponential($\theta=0.57735$)+(a, $\Delta\mu$), n=7
 $a=-0.5371$, $\Delta\mu=-0.5774$



```
In [87]: 1 plot_pvals_with_perm()
2     distx = Uniform(-1, 1), disty = Exponential(0.5773502691896257),
3     m = 5, n = 10, L = 10^4)
```

3.106114 seconds (10.20 k allocations: 1.440 MiB)
0.001654 seconds (138 allocations: 249.516 KiB)
0.001222 seconds (122 allocations: 246.500 KiB)
 $a = -0.5370568188698568$
 $\Delta\mu = -0.5773502691896257$

Out[87]: X: Uniform(a=-1.0, b=1.0), m=5
Y: Exponential($\theta=0.57735$)+(a, $\Delta\mu$), n=10
 $a=-0.5371$, $\Delta\mu=-0.5774$



In [88]:

```
1 plot_pvals_with_perm()
2     distx = Uniform(-1, 1), disty = Exponential(0.5773502691896257),
3     m = 10, n = 5, L = 10^4)
```

3.331237 seconds (10.20 k allocations: 1.897 MiB)

0.001801 seconds (141 allocations: 249.609 KiB)

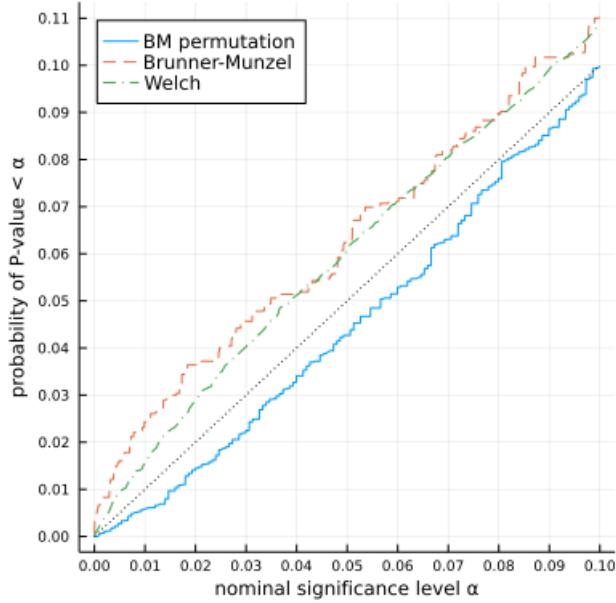
0.001320 seconds (118 allocations: 246.375 KiB)

a = -0.5370568188698568

$\Delta\mu = -0.5773502691896257$

Out[88]:

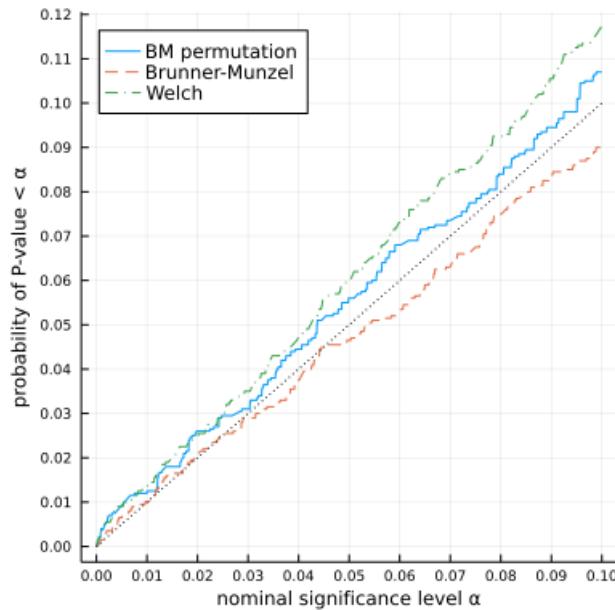
X: Uniform(a=-1.0, b=1.0), m=10
Y: Exponential($\theta=0.57735$) $+(a, \Delta\mu)$, n=5
 $a=-0.5371, \Delta\mu=-0.5774$



```
In [89]: 1 plot_pvals_with_perm()
2     distx = Uniform(-1, 1), disty = Exponential(0.5773502691896257),
3     m = 10, n = 10, L = 2000)
```

47.644018 seconds (2.26 k allocations: 17.259 MiB)
 0.000470 seconds (132 allocations: 63.281 KiB)
 0.025918 seconds (115 allocations: 59.641 KiB)
 $a = -0.5370568188698568$
 $\Delta\mu = -0.5773502691896257$

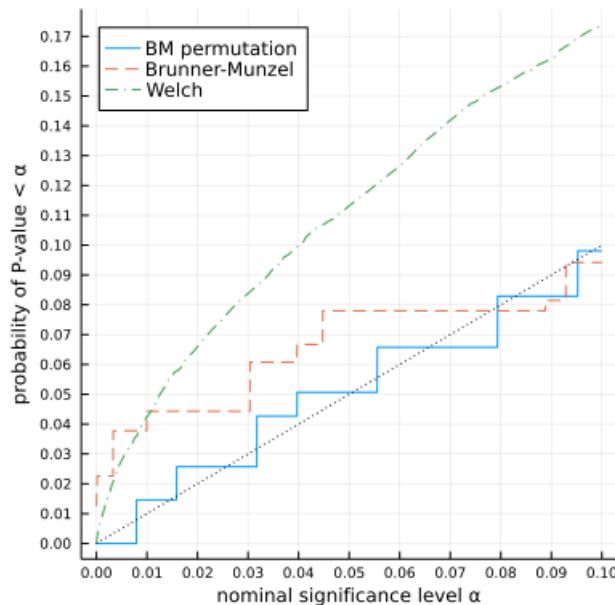
Out[89]: X: Uniform(a=-1.0, b=1.0), m=10
 Y: Exponential($\theta=0.57735$)+(a, $\Delta\mu$), n=10
 $a=-0.5371$, $\Delta\mu=-0.5774$



```
In [90]: 1 plot_pvals_with_perm()
2     distx = LogNormal(), disty = LogNormal(1), m = 5, n = 5, L = 10^4)
```

0.251410 seconds (35.39 k allocations: 2.456 MiB, 15.98% compilation time)
 0.001826 seconds (146 allocations: 249.109 KiB)
 0.001478 seconds (112 allocations: 245.859 KiB)
 $a = -1.4744426128871542$
 $\Delta\mu = -2.8329677996379363$

Out[90]: X: LogNormal($\mu=0.0$, $\sigma=1.0$), m=5
 Y: LogNormal($\mu=1.0$, $\sigma=1.0$)+(a, $\Delta\mu$), n=5
 $a=-1.4744$, $\Delta\mu=-2.833$



In [91]:

```

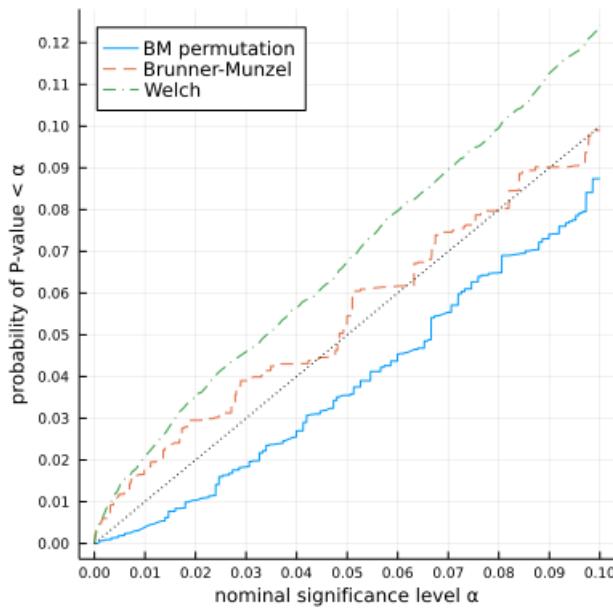
1 plot_pvals_with_perm(
2     distx = LogNormal(), disty = LogNormal(1), m = 5, n = 10, L = 10^4)

3.130992 seconds (10.19 k allocations: 1.440 MiB)
0.001864 seconds (243 allocations: 258.625 KiB)
0.002055 seconds (112 allocations: 246.422 KiB)
a = -1.4744426128871542
Δμ = -2.8329677996379363

```

Out[91]:

X: $\text{LogNormal}(\mu=0.0, \sigma=1.0)$, m=5
Y: $\text{LogNormal}(\mu=1.0, \sigma=1.0)+(a, \Delta\mu)$, n=10
a=-1.4744, Δμ=-2.833



In [92]:

```

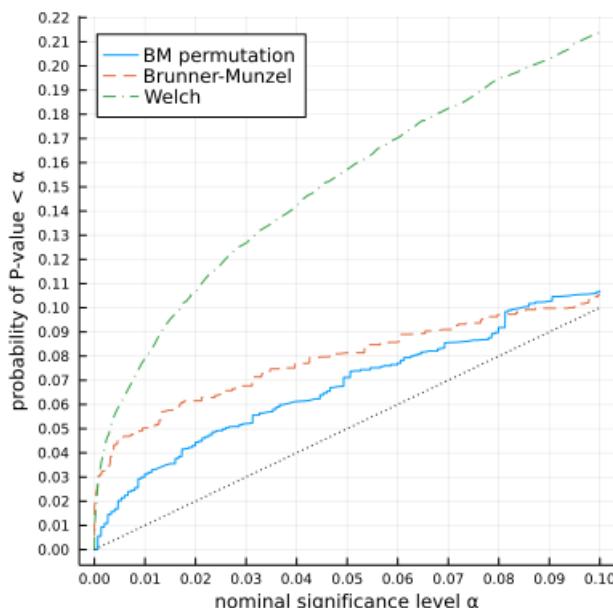
1 plot_pvals_with_perm(
2     distx = LogNormal(), disty = LogNormal(1), m = 10, n = 5, L = 10^4)

3.197243 seconds (10.19 k allocations: 1.897 MiB)
0.001818 seconds (137 allocations: 249.875 KiB)
0.001960 seconds (119 allocations: 246.594 KiB)
a = -1.4744426128871542
Δμ = -2.8329677996379363

```

Out[92]:

X: $\text{LogNormal}(\mu=0.0, \sigma=1.0)$, m=10
Y: $\text{LogNormal}(\mu=1.0, \sigma=1.0)+(a, \Delta\mu)$, n=5
a=-1.4744, Δμ=-2.833



```
In [93]: 1 plot_pvals_with_perm()  
2     distx = LogNormal(0), disty = LogNormal(1), m = 10, n = 10, L = 2000)
```

48.697102 seconds (2.27 k allocations: 17.259 MiB, 0.04% gc time)

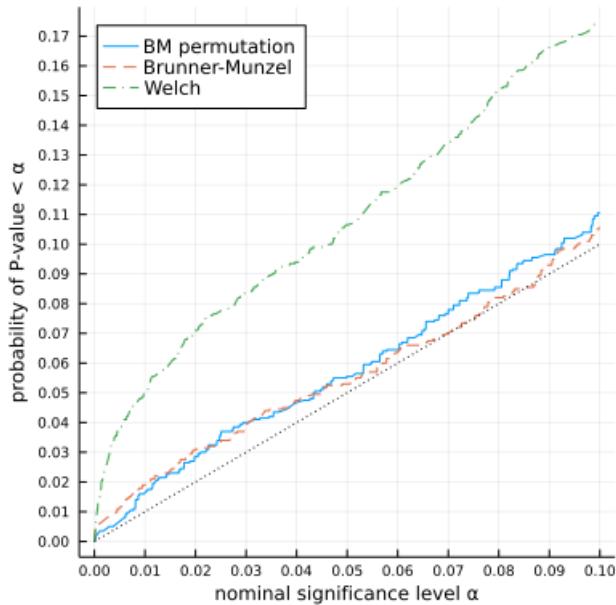
0.000675 seconds (141 allocations: 64.031 KiB)

0.001017 seconds (115 allocations: 59.859 KiB)

a = -1.4744426128871542

$\Delta\mu$ = -2.8329677996379363

```
Out[93]: X: LogNormal(mu=0.0, sigma=1.0), m=10  
Y: LogNormal(mu=1.0, sigma=1.0)+(a, Delta mu), n=10  
a=-1.4744, Delta mu=-2.833
```



```
In [ ]: 1
```