

Programming Updates in Maple 2023

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ArrayTools

```
> with( ArrayTools );
```

SortBy

- The new [ArrayTools:-SortBy](#) command allows one to easily sort a two-dimensional Array (like a Matrix) by a specific column or row. For example:

```
> A := Matrix( [[5,1,3,8,1],[20,18,12,0,22],[15,3,11,15,3],[4,10,19,3,22]] );
```

$$A := \begin{bmatrix} 5 & 1 & 3 & 8 & 1 \\ 20 & 18 & 12 & 0 & 22 \\ 15 & 3 & 11 & 15 & 3 \\ 4 & 10 & 19 & 3 & 22 \end{bmatrix}$$

```
> B := SortBy( A, 'column', 3 );
```

$$B := \begin{bmatrix} 5 & 1 & 3 & 8 & 1 \\ 15 & 3 & 11 & 15 & 3 \\ 20 & 18 & 12 & 0 & 22 \\ 4 & 10 & 19 & 3 & 22 \end{bmatrix}$$

- Custom sorting options and keywords are supported, as is in-place sorting:

```
> SortBy( A, 'row', 2, 'descending', 'inplace' );
```

```
> 'A' = A;
```

$$A = \begin{bmatrix} 1 & 5 & 1 & 3 & 8 \\ 22 & 20 & 18 & 12 & 0 \\ 3 & 15 & 3 & 11 & 15 \\ 22 & 4 & 10 & 19 & 3 \end{bmatrix}$$

IsSubsequence

- The new [ArrayTools:-IsSubsequence](#) command checks if a one-dimensional container is a subsequence of another. More precisely, **A** is a subsequence of **B** when there is a list of strictly increasing indices **K** such that **A=B[K]**. For example:

```
> IsSubsequence( <1,3,5>, <1,2,3,4,5> );
                                         true
> IsSubsequence( <1,5,3>, <1,2,3,4,5> );
                                         false
> IsSubsequence( <2,4,6>, <1,2,3,4,5> );
                                         false
```

- The **match** option is used to customize how matches are determined. For example, to check a subsequence of floats, one can do the following:

```
> B := Array( [ exp(1), Pi, gamma ] );
                                         B := [ e π γ ]
> A := evalf[5]( B[[2,3]] );
                                         A := [ 3.1416 0.57722 ]
> IsSubsequence( A, B );
                                         false
> IsSubsequence( A, B, 'match' = 'float' );
                                         false
> IsSubsequence( A, B, 'match' = 'float', 'digits' = 5, 'ulp' = 1 )
;
                                         true
```

- The matching indices can also be returned:

```
> A := [ "a", "e", "i", "o", "u" ];
                                         A := [ "a", "e", "i", "o", "u" ]
> B := [ seq( "a" .. "z" ) ];
                                         B := [ "a", "b", "c", "d", "e", "f", "g", "h", "i", "j", "k", "l", "m", "n", "o", "p", "q", "r", "s", "t", "u", "v",
                                         "w", "x", "y", "z" ]
> K := IsSubsequence( A, B, 'output' = 'indices' );
                                         K := [ 1, 5, 9, 15, 21 ]
> EqualEntries( A, B[K] );
                                         true
```

ColorTools

- The [ColorTools](#) package now supports the CAM02 color spaces **JCh** (cylindrical) and **Jab** (rectangular). These color spaces are now the most modern perceptual color spaces in Maple. JCh is useful for making perceptually uniform changes to lightness (J), chromaticity/saturation (C), and hue (h).

```
> ColorTools:-Color("JCh", "#9f1");  
                                ⟨JCh : 0.852 0.926 0.353⟩
```

- Jab is useful as a space where Euclidean distance is roughly the same as perceptual distance. These different applications are the reason why the **J** channel is not the same in the two spaces. [ColorTools:-Gradient](#) now uses **Jab** for its **best** mode.

```
> ColorTools:-Color("Jab", "#9f1");  
                                ⟨Jab : 0.907 -0.245 0.324⟩
```

- The **YUV** rectangular color space is now supported, using exactly the same implementation as in [ImageTools:-RGBtoYUV](#) and [ImageTools:-YUVtoRGB](#). The **Y** channel is lightness, while **U** and **V** are rectangular coordinates of chromacity.

```
> ColorTools:-Color("YUV", "#9f1");  
                                ⟨YUV : 0.774 -0.348 -0.153⟩
```

- The **HSL** cylindrical color space is now supported using a simple conversion from the existing **HSV** color space. The **H** is always the same between the two but the **S** saturation channel differs.

```
> ColorTools:-Color("HSL", "#9f1");  
                                ⟨HSL : 0.238 1 0.533⟩
```

```
> ColorTools:-Color("HSV", "#9f1");  
                                ⟨HSV : 0.238 0.933 1⟩
```

- A new grayscale color space is now supported called **Gs**. Colors are converted to grayscale by calculating their perceptual lightness.

```
> ColorTools:-Color("Gs", "Red");  
                                ⟨Gs : 0.532⟩
```

- Colors do not roundtrip convert due to loss of information about hue and chroma but all true shades of gray do.

```
> ColorTools:-Color("RGB", ColorTools:-Color("Gs", "#7e7e7e")) =  
ColorTools:-Color("RGB", "#7e7e7e");  
                                ⟨RGB : 0.494 0.494 0.494⟩ = ⟨RGB : 0.494 0.494 0.494⟩
```

- A new conversion **color** has been added to [convert](#) so that [convert/color](#) can be used to convert easily between supported color formats.

```
> convert( ColorTools:-Color("Jab", "#9f1"), 'color', "hex");
          "#99FF11"
```

and between color spaces

```
> convert( ColorTools:-Color("Jab", "#9f1"), 'color', "HSL");
          ⟨HSL : 0.238 1 0.533⟩
```

```
> convert( [0.6, 1.0, 0.066666667], 'color', "RGB", "YUV");
          [0.774000000038000, -0.348133333188000, -0.152666666700000]
```

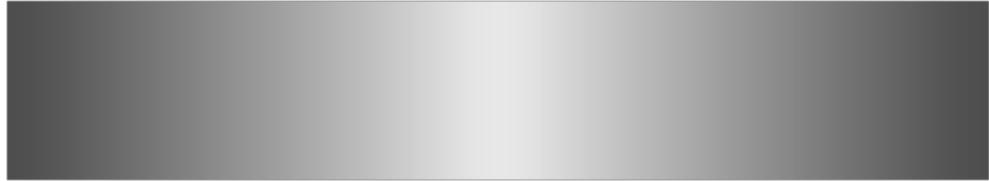
```
> convert( [0, 128, 0], 'color', "colorname");
          "Green"
```

- The [ColorTools:-Swatches](#) command now has two new options **mode** and **filter**. The mode option applies one of several preset styles. The filter option applies a procedure to each color before displaying it.

```
> ColorTools:-Swatches("Colorwheel", 'mode'='wheel','filter'=
(c->ColorTools:-CVDSimulation(c,"tritanomaly")));
```



```
> ColorTools:-Swatches("Colorwheel", 'mode'='gradient', 'filter'=
(c->ColorTools:-Color("Gs",c)));
```



ListTools

- Two new related commands, [ListTools:-InversePermutation](#) and [ListTools:-Unpermute](#), determine, respectively, the inverse of a permutation given the forward permutation, and an unpermuted list given the permuted list and forward permutation. For example:

```
> with( ListTools ):

> X := [seq("a".."j")]; # unpermuted list
X := ["a", "b", "c", "d", "e", "f", "g", "h", "i", "j"]

> P := [4,6,3,9,8,10,1,7,2,5]; # forward permutation
P := [4, 6, 3, 9, 8, 10, 1, 7, 2, 5]

> Y := X[P]; # permuted list
Y := ["d", "f", "c", "i", "h", "j", "a", "g", "b", "e"]

> Q := InversePermutation(P); # inverse permutation
Q := [7, 9, 3, 1, 10, 2, 8, 5, 4, 6]

> Y[Q]; # unpermuted list
["a", "b", "c", "d", "e", "f", "g", "h", "i", "j"]

> Unpermute(Y,P); # unpermuted list
["a", "b", "c", "d", "e", "f", "g", "h", "i", "j"]
```

membertype

- The **membertype(T,S)** function determines if there is an operand within the expression S of type T. In Maple 2023 [membertype](#) has been extended to support a wider variety of base expressions, S, to scan. It now supports products, sums, and rtables. For example:

```
> membertype( integer, 97*x+4 );
true

> membertype( integer, Matrix([[1,2],[3,4]]) );
true
```

New inverse functions for Box objects

- For Maple 2023, **RealBox** and **ComplexBox** objects now support the inverse circular functions **arcsec**, **arccsc**, and **arccot**, as well as the inverse hyperbolic functions **arcsech**, **arccsch**, and **arccoth**.

```
> arcsec( RealBox( 4.7 ) );
                                         (RealBox: 1.35639 ± 2.7043e-10)

> arcsech( RealBox( 0.23 ) );
                                         (RealBox: 2.14933 ± 4.1295e-10)

> arccot( ComplexBox( 4.3*I - 0.33 ) );
                                         (ComplexBox: [3.12285 +/- 4.38103e-10] + [-0.235372 +/- 2.71693e-10]*I)
```

- For more information, see [RealBox](#) and [ComplexBox](#).

New Units prefixes

- In November, 2022, the 27th meeting of the General Conference on Weights and Measures approved the new prefixes quetta-, ronna-, ronto-, and quecto- for SI units, meaning factors of 10^{30} , 10^{27} , 10^{-30} , and 10^{-27} , respectively. These prefixes are fully integrated into Maple 2023. We can express the Earth's mass in ronnagrams and the Sun's in quettagrams.

```
> restart;

> with(ScientificConstants):

> m_earth := Constant(M[Earth], units):

> m_earth := convert(evalf(m_earth), units, ronnagrams);
                                         m_earth := 5.972190000 Rg

> m_sun := Constant(M[Sun], units):

> m_sun := convert(evalf(m_sun), units, Qg);
                                         m_sun := 1990.000000 Qg
```

- The average (Newtonian) gravitational force between the Earth and the Sun is most compactly expressed in zettanewtons. (The prefix zetta-, meaning a factor of 10^{21} , was introduced in 1991.)

```
> G := evalf(Constant(G, units));
                                         3
                                         G := 6.67408 × 10  −11  m
                                         kg s 2
```

```

> force_earth_sun := convert(G * m_sun * m_earth / Unit(AU^2),
  units, zettanewton);
  forceearth_sun := 35.44273899 ZN

• The typical (Newtonian) gravitational force between two cups of tea, one on Earth and
one on the moon, can be computed as follows.

> volume_tea := Unit(cup);
  volumetea := cup

> density_tea := ThermophysicalData:-Property("density", "water", T=
95*Unit(Celsius), pressure=Unit(atm));
  densitytea := 961.8879166 kg/m3

> mass_tea := volume_tea * density_tea;
  masstea := 961.8879166 cup kg/m3

> distance_moon := 384400*Unit(km);
  distancemoon := 384400 km

> force_tea_tea := convert(G * mass_tea^2 / distance_moon^2, units,
qN);
  forcetea_tea := 23.39159750 qN

• We can express the Planck length in quectometers.

> planck := Constant(l[P], units):
> convert(evalf(planck), units, quectometer);
  0.00001616228373 qm

```